

Figure 1S

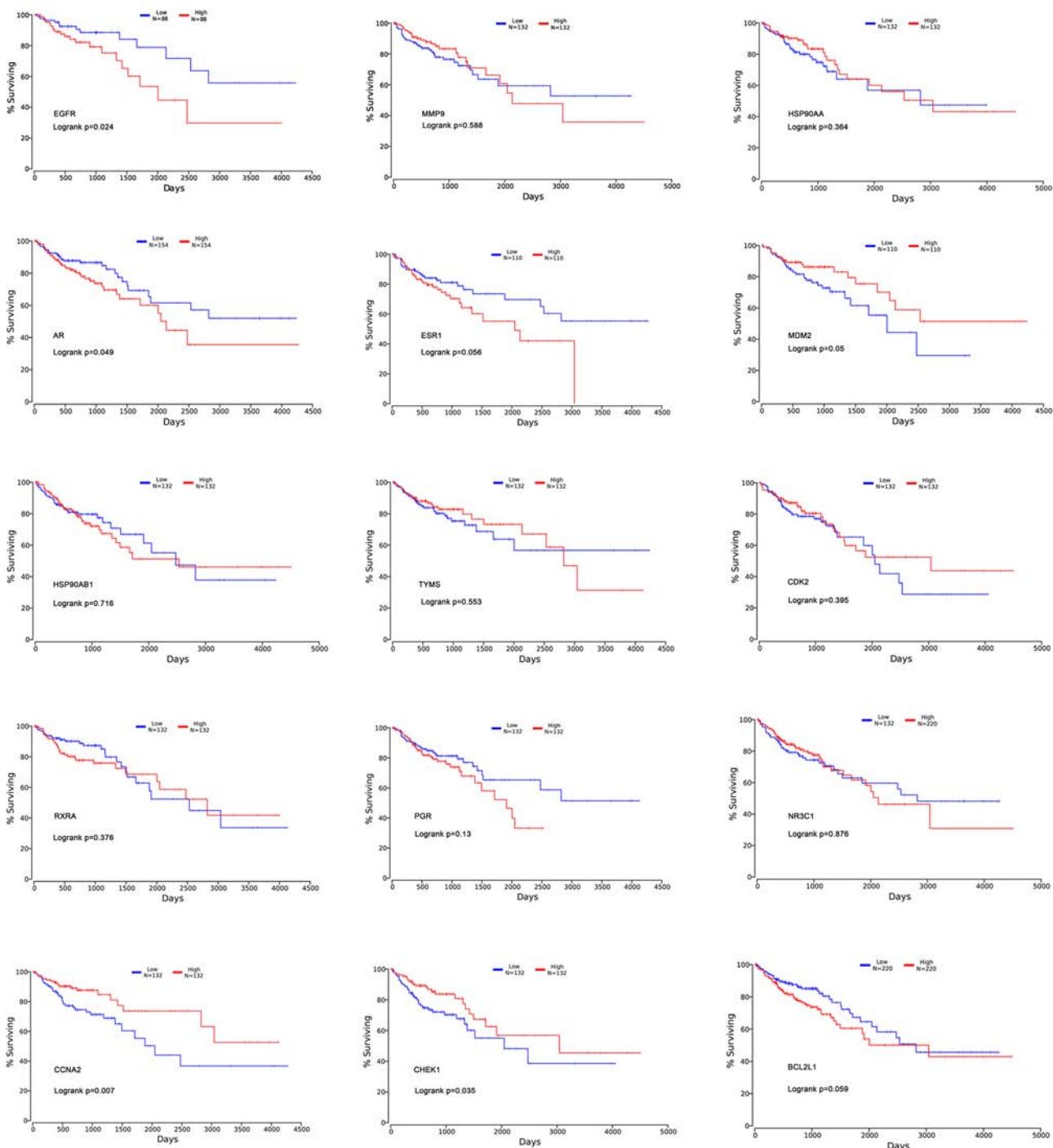


Table 1S

| Statistics of MTS experiments of SW480 and RKO | | | | | | | | | |
|------------------------------------------------|---|-------------------------------|---------|---------|---------|-----------------------------|---------|---------|---------|
| | | SW480 (Compared with 0 μ mol) | | | | RKO (Compared with 0 μ mol) | | | |
| Time | | 1 | 5 | 10 | 20 | 1 | 5 | 10 | 20 |
| 24h | t | 2.133 | 2.115 | 5.496 | 5.538 | 0.8428 | 1.169 | 1.832 | 1.614 |
| | p | 0.0769 | 0.0788 | 0.0015 | 0.0015 | 0.4315 | 0.2868 | 0.1167 | 0.1576 |
| 48h | t | 0.7661 | 9.239 | 15.24 | 28.72 | 0.0354 | 3.13 | 5.1 | 5.718 |
| | p | 0.4727 | <0.0001 | <0.0001 | <0.0001 | 0.9729 | 0.0203 | 0.0022 | 0.0012 |
| 72h | t | 0.7101 | 8.088 | 14.36 | 19.06 | 1.079 | 19.83 | 14.57 | 54.43 |
| | p | 0.5043 | 0.0002 | <0.0001 | <0.0001 | 0.3222 | <0.0001 | <0.0001 | <0.0001 |
| 96h | t | 3.514 | 19.3 | 16.05 | 27.52 | 1.518 | 16.52 | 40.25 | 49.83 |
| | p | 0.0126 | <0.0001 | <0.0001 | <0.0001 | 0.1798 | <0.0001 | <0.0001 | <0.0001 |

Ligand:
Table 2S 10322911

| | | | | | | | | | | | | | | | |
|----------|---------|---|-------|--------|------------|---|---|---|---|---|----------------------------------------------------------|-----------------|---------------|---------------|------|
| CSNK2A1 | 3h30_v | 4 | 2.437 | 0.6092 | 0.0584914 | 2 | 0 | 2 | 0 | 0 | 0 Casein kiNONE | P68400 | Involved NONE | Being inv | |
| HSPA8 | 3fzk_v | 4 | 2.436 | 0.6091 | -0.253067 | 2 | 2 | 0 | 0 | 0 | 0 Heat shocNONE | P11142 | MolecularNONE | NONE | |
| TTPA | 1oiz_v | 5 | 3.044 | 0.6088 | 0.666902 | 4 | 1 | 0 | 0 | 0 | 0 Alpha-tocNONE | P49638 | Involved NONE | Vitamin E | |
| PYGL | 1fa9_v | 4 | 2.432 | 0.608 | -0.287533 | 0 | 2 | 2 | 0 | 0 | 0 Glycogen NONE | P06737 | CarbohydrNONE | For nutri | |
| MMP3 | 1d7x_v | 4 | 2.424 | 0.6061 | -0.0557303 | 1 | 1 | 2 | 0 | 0 | 0 StromelysHYDROLASEMMP3_HUMACan degraNONE | | | NONE | |
| GSK3B | 3f7z_v | 5 | 3.029 | 0.6058 | 0.913089 | 3 | 2 | 0 | 0 | 0 | 0 Glycogen NONE | P49841 | Involved NONE | NONE | |
| HDAC8 | 1w22_v | 6 | 3.603 | 0.6005 | 1.6543 | 3 | 1 | 2 | 0 | 0 | 0 Histone dHYDROLASEHDAC8_HUMResponsibNONE | | | NONE | |
| DPP4 | 2hha_v | 5 | 2.992 | 0.5984 | 1.11623 | 2 | 2 | 0 | 1 | 0 | 0 DipeptidylHYDROLASEDPP4_HUMARemoves NNONE | | | Rheumatoi | |
| PNMT | 1n7i_v | 4 | 2.39 | 0.5976 | -0.86091 | 3 | 0 | 0 | 1 | 0 | 0 PhenylethTRANSFERNAMT_HUMAConverts NNONE | | | NONE | |
| DHDH | 1d3g_v | 4 | 2.384 | 0.5961 | -0.6868 | 3 | 0 | 1 | 0 | 0 | 0 DihydrodroOXIDOREDUYRD_HUMANONE | | | NONE | |
| LCK | 2of4_v | 4 | 2.384 | 0.5961 | -0.234403 | 2 | 1 | 0 | 1 | 0 | 0 Proto-oncTRANSFERALCK_HUMANtyrosine A chromosPsoriasis | | | | |
| PDE3B | 1so2_v | 5 | 2.954 | 0.5908 | 0.797327 | 2 | 2 | 1 | 0 | 0 | 0 cGMP-inhiHYDROLASEPDE3B_HUMMay play NONE | | | NONE | |
| CFD | 1dic_v | 4 | 2.352 | 0.5881 | -0.614332 | 2 | 2 | 0 | 0 | 0 | 0 ComplementSERINE_PRCFAD_HUMAFactor D Defects | iAutoimmun | | | |
| CTSG | 1chg_v | 5 | 2.935 | 0.587 | 0.975307 | 1 | 2 | 0 | 2 | 0 | 0 CathepsinCOMPLEX (CATG_HUMASerine prNONE | Connectiv | | | |
| PARP1 | 1wok_v | 5 | 2.934 | 0.5867 | 0.494421 | 3 | 1 | 1 | 0 | 0 | 0 Poly [ADPNONE | P09874 | Involved NONE | NONE | |
| PGF | 1fvz_v | 5 | 2.931 | 0.5862 | 0.720473 | 3 | 0 | 2 | 0 | 0 | 0 Placenta HORMONE/GPLGF_HUMAGrowth faNONE | Retinopat | | | |
| CPB1 | 1kwm_v | 5 | 2.927 | 0.5855 | -0.10423 | 0 | 3 | 0 | 0 | 2 | 0 | 0 CarboxypeNONE | P15086 | Involved NONE | NONE |
| AMY2A | 3cpu_v | 5 | 2.918 | 0.5837 | 0.907559 | 0 | 2 | 3 | 0 | 0 | 0 PancreatiHYDROLASEAMY_HUMANONE | | | Bacillus | |
| F2 | 1ktt_v | 5 | 2.905 | 0.5811 | 0.571482 | 3 | 1 | 0 | 1 | 0 | 0 ProthrombHYDROLASETHRB_HUMATHrombin,Defects | iDeep vein | | | |
| ISG20 | 1wlj_v | 5 | 2.899 | 0.5798 | 0.255112 | 0 | 4 | 1 | 0 | 0 | 0 InterferoHYDROLASEISG20_HUMExonucleaNONE | | | NONE | |
| LTAAH | 3cho_v | 4 | 2.307 | 0.5767 | -0.149397 | 2 | 1 | 0 | 1 | 0 | 0 LeukotriehHYDROLASELKA_HUMHydrolizeNONE | Inflammatt | | | |
| PDE5A | 1rkp_v | 4 | 2.296 | 0.5741 | -0.424825 | 1 | 2 | 0 | 0 | 0 | 1 cGMP-spECHYDROLASEPDE5A_HUMPlays a rNONE | Anal fiss | | | |
| ITK | 1sm2_v | 6 | 3.437 | 0.5729 | 1.3578 | 4 | 2 | 0 | 0 | 0 | 0 Tyrosine-NONE | Q08881 | Involved NONE | NONE | |
| ADK | 2i6b_v | 6 | 3.421 | 0.5702 | 1.04405 | 5 | 1 | 0 | 0 | 0 | 0 AdenosineNONE | P55263 | ATP depenNONE | NONE | |
| PLAU | 1f92_v | 5 | 2.847 | 0.5695 | 0.756844 | 1 | 2 | 1 | 1 | 0 | 0 UrokinaseHYDROLASEUROK_HUMASpecificaNONE | Angiogene | | | |
| MMP12 | 1utt_v | 5 | 2.846 | 0.5693 | 0.373809 | 2 | 3 | 0 | 0 | 0 | 0 MacrophagHYDROLASEMMP12_HUMMay be irNONE | Chronic o | | | |
| HMGCR | 2q6c_v | 5 | 2.845 | 0.5691 | 0.516261 | 2 | 2 | 0 | 0 | 1 | 0 3-hydroxyHYDROREDUHMDH_HUMATHis tranNONE | Alzheimer | | | |
| CYP2C9 | log5_v | 6 | 3.414 | 0.5689 | 1.61603 | 3 | 3 | 0 | 0 | 0 | 0 CytochromELECTRON CP2C9_HUMCytochromNONE | | | NONE | |
| CBR1 | 1wma_v | 5 | 2.834 | 0.5668 | 0.502088 | 2 | 2 | 1 | 0 | 0 | 0 Carbonyl OXIDOREDUCBR1_HUMACatalyzesNONE | | | NONE | |
| EPHX2 | 1vjt5_v | 5 | 2.818 | 0.5636 | 0.181816 | 3 | 1 | 1 | 0 | 0 | 0 Epoxide hHYDROLASEHYES_HUMAActs on eNONE | | | NONE | |
| ADH1C | 1u3w_v | 5 | 2.812 | 0.5625 | 0.25127 | 3 | 1 | 0 | 1 | 0 | 0 Alcohol dNONE | P00326 | Energy prNONE | Some evid | |
| AHCY | 1a7a_v | 5 | 2.809 | 0.5617 | 0.123104 | 0 | 3 | 2 | 0 | 0 | 0 AdenosylHYDROLASESAHH_HUMAAdenosylhDefects | iInflammatt | | | |
| ALDH2 | 1of7_v | 5 | 2.806 | 0.5612 | 0.603611 | 2 | 1 | 2 | 0 | 0 | 0 Aldehyde OXIDOREDUALDH2_HUNNONE | | | NONE | |
| SERPINA1 | 1nz2_v | 6 | 3.359 | 0.5598 | 0.55942 | 3 | 1 | 2 | 0 | 0 | 0 Alpha-1-aPROTEINase inhibitorThe major Psoriasis | | | | |
| EPHB4 | 2vww_v | 4 | 2.238 | 0.5595 | -0.727475 | 3 | 0 | 1 | 0 | 0 | 0 Ephrin tyTRANSFERAEPHB4_HUMReceptor | Ocular di | | | |
| FGFR1 | 1agw_v | 4 | 2.219 | 0.5549 | -0.81237 | 2 | 1 | 1 | 0 | 0 | 0 Basic filoKFGFR1_HUMReceptor Defects | iNONE | | | |
| JAK3 | 1yv_j_v | 6 | 3.327 | 0.5545 | 1.19431 | 4 | 1 | 1 | 0 | 0 | 0 Tyrosine-TRANSFERAJAK3_HUMATyrosine Defects | iPsoriasis | | | |
| CTNN1A | 1h6g_v | 5 | 2.771 | 0.5541 | 0.240449 | 3 | 2 | 0 | 0 | 0 | 0 Catenin aCYTOSKELETNAL_HUMAssociateAbnormali | | | | |
| WAS | 1t84_v | 5 | 2.759 | 0.5518 | -0.382247 | 5 | 0 | 0 | 0 | 0 | 0 Wiskott-A-SIGNALINGWASP_HUMAEffector Defects | iWiskott-A | | | |
| TRAPPC3 | 1sz7_v | 6 | 3.298 | 0.5496 | 0.894174 | 5 | 1 | 0 | 0 | 0 | 0 TraffickiNONE | Q43617 | May play NONE | NONE | |
| SOD2 | 2ixil_v | 5 | 2.748 | 0.5496 | 0.276764 | 2 | 2 | 1 | 0 | 0 | 0 SuperoxidNONE | P04179 | Destroys NONE | NONE | |
| GLRX | 1b4q_v | 5 | 2.733 | 0.5466 | 0.21955 | 1 | 1 | 0 | 1 | 2 | 0 GlutaredoxOXYDREDUGLRX1_HUMHas gluNONE | | | NONE | |
| FABP7 | 1fe3_v | 7 | 3.826 | 0.5465 | 1.33678 | 6 | 0 | 0 | 0 | 1 | 0 Fatty aciLIPID BINFABP7_HUMB-FABP ccNONE | | | NONE | |
| REG1A | 1qdd_v | 5 | 2.725 | 0.545 | 0.392411 | 0 | 4 | 1 | 0 | 0 | 0 LithostatNONE | P05451 | Involved NONE | NONE | |
| NR1H2 | 1upw_v | 6 | 3.266 | 0.5444 | 0.979271 | 5 | 0 | 1 | 0 | 0 | 0 OxysterolRECEPTOR NR1H2_HUMOrphan reNONE | Lipid met | | | |
| FGFR2 | 1oec_v | 5 | 2.703 | 0.5406 | -0.243324 | 3 | 1 | 0 | 1 | 0 | 0 FibroblastNONE | P21802 | Involved NONE | For treat | |
| NR1H3 | 1uh1_v | 6 | 3.243 | 0.5405 | 1.01947 | 5 | 1 | 0 | 0 | 0 | 0 OxysterolDNA BINDINR1H3_HUMOrphan reNONE | | | NONE | |
| PPRA | 1i7g_v | 6 | 3.233 | 0.5388 | 1.10437 | 4 | 2 | 0 | 0 | 0 | 0 PeroxisomTRANSCRIPPPARA_HUMReceptor | Syndrome | | | |
| MTHFD1 | 1dia_v | 6 | 3.232 | 0.5387 | 1.69957 | 2 | 2 | 2 | 0 | 0 | 0 C-1-tetraNONE | P11586 | NucleotidNONE | For nutri | |
| NR1H4 | 1osh_v | 5 | 2.681 | 0.5362 | -0.314068 | 5 | 0 | 0 | 0 | 0 | 0 Bile acid TRANSCRIPNR1H4_HUMReceptor | | | Liver fib | |
| #N/A | 1ydt_v | 5 | 2.671 | 0.5343 | -0.254979 | 4 | 0 | 0 | 1 | 0 | 0 cAMP-depeNONE | P00517 | Involved NONE | NONE | |
| AKR1C1 | 1mrq_v | 5 | 2.656 | 0.5311 | -0.19207 | 4 | 1 | 0 | 0 | 0 | 0 Aldo-ketoNONE | P04828 | Involved NONE | Some evid | |
| #N/A | 2vd1_v | 6 | 3.181 | 0.5301 | 0.566682 | 5 | 1 | 0 | 0 | 0 | 0 GlutathioISOMERASEPTG2_HUMCatalyzesNONE | | | NONE | |
| GL01 | 1qip_v | 7 | 3.71 | 0.53 | 1.05243 | 3 | 1 | 0 | 1 | 2 | 0 LactoylglyLYASE LGUL_HUMACatalyzesNONE | Cancer | | | |
| SULT1E1 | 1g3m_v | 7 | 3.708 | 0.5297 | 1.4124 | 6 | 1 | 0 | 0 | 0 | 0 Estrogen TRANSFERAST1E1_HUMMay contrNONE | | | NONE | |
| THRA | 1nav_v | 7 | 3.68 | 0.5257 | 1.49423 | 4 | 2 | 1 | 0 | 0 | 0 Thyroid hNONE | P10827 | Involved NONE | Used as r | |
| SYK | 1xbc_v | 5 | 2.615 | 0.523 | -0.274543 | 3 | 1 | 1 | 0 | 0 | 0 Tyrosine-TRANSFERAKSYK_HUMAPositive NONE | Systemic | | | |
| #N/A | 1gzz_v | 6 | 3.129 | 0.5215 | 1.04478 | 3 | 2 | 0 | 1 | 0 | 0 Insulin-1NONE | P01343 | NONE | NONE | |
| ADAM17 | 3edz_v | 5 | 2.597 | 0.5194 | 0.413479 | 2 | 2 | 0 | 0 | 0 | 1 DisintegrNONE | P78536 | Cleaves tNONE | NONE | |
| RBP4 | 1rbp_v | 8 | 4.134 | 0.5168 | 1.03618 | 8 | 0 | 0 | 0 | 0 | 0 Retinol-hRETINOL IRET4_HUMDelivers | iNONE | | | |
| GSTT2B | 1ljr_v | 7 | 3.601 | 0.5144 | 2.02054 | 1 | 3 | 1 | 1 | 1 | 0 GlutathioTRANSFERTAGSTT2_HUMConjugatiNONE | | | NONE | |
| STGA1 | 1gs_e_v | 6 | 3.085 | 0.5142 | 0.400347 | 5 | 1 | 0 | 0 | 0 | 0 GlutathioNONE | P08263 | Involved NONE | For nutri | |
| XIAP | 2opy_v | 5 | 2.561 | 0.5123 | 0.124794 | 1 | 2 | 0 | 1 | 0 | 0 BaculovirAPOPTOSISXIAP_HUMApoptoticDefects | iCancer;Sk | | | |
| FABP6 | 1olv_v | 7 | 3.58 | 0.5114 | 1.04953 | 5 | 1 | 1 | 0 | 0 | 0 GastrotroLIPID BINFABP6_HUMIleal prNONE | | | NONE | |
| FABP3 | 2hmb_v | 7 | 3.559 | 0.5084 | 0.878363 | 5 | 1 | 0 | 0 | 1 | 0 Fatty aciLIPID-BINFABPH_HUMFABP are NONE | | | NONE | |
| PLA2G2A | 1dcy_v | 5 | 2.528 | 0.5056 | -0.954748 | 3 | 1 | 0 | 0 | 1 | 0 PhospholipNONE | P14555 | Involved NONE | NONE | |
| CSNK1G2 | 2c47_v | 4 | 2.02 | 0.5051 | -1.20191 | 1 | 1 | 2 | 0 | 0 | 0 Casein kiTRANSFERAKC1G2_HUMCasein kiNONE | | | NONE | |
| YARS | 1q11_v | 5 | 2.525 | 0.5049 | -0.40966 | 2 | 1 | 1 | 1 | 0 | 0 Tyrosyl-tNONE | P54577 | TranslatiNONE | Tyrosine | |
| CDK6 | 1x02_v | 5 | 2.523 | 0.5047 | -0.18693 | 2 | 3 | 0 | 0 | 0 | 0 Cell diviCELL CYCLCDK6_HUMAProbably NONE | | | NONE | |
| RNASE4 | 2rnf_v | 5 | 2.507 | 0.5013 | -0.462335 | 0 | 3 | 1 | 0 | 1 | 0 RibonucleNONE | P34096 | Involved NONE | NONE | |
| FHIT | 2fhi_v | 5 | 2.502 | 0.5004 | 0.0817782 | 0 | 3 | 1 | 0 | 0 | 1 Bis(5-adeNONE | P49789 | NucleotidNONE | NONE | |
| MET | 1r0p_v | 6 | 2.975 | 0.4958 | 0.121518 | 4 | 2 | 0 | 0 | 0 | 0 Hepatocyt TRANSFERAMET_HUMANReceptor Activatio | Alopecia, | | | |
| PDK2 | 2bu5_v | 6 | 2.967 | 0.4945 | 0.0375573 | 5 | 1 | 0 | 0 | 0 | 0 PyruvateTRANSFERPDK2_HUMAInhibits NONE | | | NONE | |
| NR113 | 1xvp_v | 6 | 2.966 | 0.4943 | 0.241254 | 6 | 0 | 0 | 0 | 0 | 0 Nuclear rDNA BINDINR113_HUMBinds andNONE | | | NONE | |
| PK1 | 1nhx_v | 6 | 2.956 | 0.4926 | 0.0905391 | 1 | 3 | 0 | 0 | 2 | 0 | 0 PhosphoenONE | P35558 | Energy prNONE | NONE |
| F7 | 2flr_v | 5 | 2.421 | 0.4842 | -0.559078 | 3 | 0 | 2 | 0 | 0 | 0 CoagulatiHYDROLASEF7_HUMANInitiates | iDeep vein | | | |
| BLVRB | 1he2_v | 6 | 2.901 | 0.4835 | 0.246847 | 4 | 1 | 0 | 0 | 1 | 0 Flavin reNONE | P30043 | Cell wallNONE | For the t | |
| RHOA | 1az2b_v | 6 | 2.894 | 0.4823 | 0.915877 | 0 | 4 | 2 | 0 | 0 | 0 TransformNONE | P61586 | Involved NONE | For the t | |
| DPEP1 | 1itu_v | 8 | 3.841 | 0.4802 | 1.27229 | 4 | 2 | 1 | 0 | 1 | 0 DipeptidaHYDROLASEDPEP1_HUMHydrolyzeNONE | | | NONE | |
| #N/A | 2f57_v | 5 | 2.393 | 0.4785 | -0.523768 | 3 | 1 | 1 | 0 | 0 | 0 Serine/thTRANSFERAPAK7_HUMAthe activNONE | | | NONE | |
| MDM2 | 1rv1_v | 5 | 2.388 | 0.4777 | -1.11008 | 4 | 0 | 1 | 0 | 0 | 0 E3 ubiquiLIGASE MDM2_HUMAInhibits Seems to | Leukemia; | | | |
| AMY1B | 1q4n_v | 6 | 2.859 | 0.4765 | 0.576842 | 2 | 1 | 2 | 1 | 0 | 0 Alpha-amylNONE | P04745 | CarbohydrNONE | NONE | |
| FABP5 | 1b56_v | 5 | 2.363 | 0.4727 | -0.99898 | 3 | 1 | 0 | 0 | 1 | 0 Fatty aciLIPID-BINFABP5_HUMHigh specNONE | | | NONE | |
| NR3C1 | 1nhz_v | 8 | 3.762 | 0.4702 | 1.03739 | 7 | 1 | 0 | 0 | 0 | 0 GlucocortHORMONE RGR_C_HUMANReceptor | Defects | iAlzheimer | | |
| VDR | 1s19_v | 8 | 3.758 | 0.4697 | 2.13507 | 5 | 2 | 1 | 0 | 0 | 0 Vitamin DGENE REGUVDR_HUMANNuclear hDefects | iPackets bo | | | |
| SELP | 1gr1_v | 6 | 2.817 | 0.4694 | 0.79 | | | | | | | | | | |

| | | | | | | | | | | | | | | | | | | | |
|----------|-----------|----|-------|--------|------------|---|---|---|---|---|---|---------|-----------|-------------------|-------------|--------------|--------------------|---------------------|------|
| APRT | lore_v | 7 | 3.158 | 0.4512 | 0.746162 | 0 | 4 | 3 | 0 | 0 | 0 | Adenine | pNONE | P07741 | Nucleotid | NONE | For nutri | | |
| FNTA | lssad_4_v | 7 | 3.149 | 0.4498 | 0.603779 | 5 | 0 | 0 | 2 | 0 | 0 | 0 | Protein | fTRANSFER | FNTA_HUM | Catalyzes | NONE | | |
| TNK2 | lu4d_v | 5 | 2.24 | 0.448 | -0.954582 | 0 | 2 | 3 | 0 | 0 | 0 | 0 | Activated | TRANSFER | AACK1_HUM | Downstrea | Cancer | | |
| CYP2C8 | 1pq2_v | 7 | 3.135 | 0.4478 | 0.435702 | 5 | 1 | 0 | 0 | 0 | 1 | 0 | Cytochrom | OXIDOREDU | CP2C8_HUM | Cytochrom | NONE | | |
| DCK | 1p62_v | 7 | 3.126 | 0.4466 | 0.858597 | 3 | 3 | 1 | 0 | 0 | 0 | 0 | Deoxycyt | iTRANSFER | ADCK_HUMAN | Required | NONE | | |
| PRKCQ | 1xjd_v | 7 | 3.12 | 0.4458 | 0.427984 | 4 | 1 | 1 | 1 | 0 | 0 | 0 | Protein | kTRANSFER | AKPT_C_HUM | This is a | Inflamm | | |
| NR112 | 2o9i_v | 7 | 3.109 | 0.4442 | 0.267555 | 5 | 2 | 0 | 0 | 0 | 0 | 0 | Nuclear | rTRANScriP | R112_HUM | Orphan | NONE | | |
| BIRC7 | 3f7h_v | 6 | 2.654 | 0.4424 | -0.030016 | 2 | 2 | 1 | 1 | 0 | 0 | 0 | Baculovir | None | Q96CA5 | Apoptotic | NONE | | |
| AMY1B | 1mfv_v | 6 | 2.652 | 0.442 | 0.0341128 | 0 | 2 | 4 | 0 | 0 | 0 | 0 | Salivary | None | P04745 | Carbohydr | NONE | | |
| RARB | 1xap_v | 9 | 3.949 | 0.4387 | 1.35272 | 7 | 1 | 0 | 0 | 0 | 1 | 0 | 0 | Retinoic | TRANSCRIP | RARB_HUM | This is a | Psoriasis | |
| RARG | 1fcx_v | 9 | 3.927 | 0.4364 | 1.97412 | 6 | 1 | 1 | 0 | 0 | 1 | 0 | 0 | Retinoic | GENE | REGURARG_HUM | This is a | Emphysema | |
| RARA | 1dkf_v | 8 | 3.488 | 0.436 | 1.01032 | 5 | 3 | 0 | 0 | 0 | 0 | 0 | 0 | Retinoic | HORMONE | GRARA_HUM | This is a | Squamous | |
| ZAP70 | lu59_v | 7 | 3.011 | 0.4302 | 0.251011 | 4 | 1 | 2 | 0 | 0 | 0 | 0 | 0 | Tyrosine | -TRANSFER | ZAP70_HUM | Plays a | iTransplan | |
| MMP2 | 1hov_v | 7 | 2.985 | 0.4264 | 0.451343 | 4 | 1 | 2 | 0 | 0 | 0 | 0 | 72 kDa | tyHYDROLASE | MMP2_HUM | In additi | Leu bondDefects i | | |
| FDPS | 1yq7_v | 8 | 3.408 | 0.426 | 0.430207 | 1 | 4 | 1 | 0 | 0 | 2 | 0 | 0 | Farnesy | TRANSFER | APPs_HUM | Key enzym | NONE | |
| IL2 | 1m48_v | 7 | 2.966 | 0.4237 | 0.290016 | 4 | 1 | 2 | 0 | 0 | 0 | 0 | 0 | Interleuk | CYTOKINE | IL2_HUMAN | Produced a | Cromos | |
| MAP2K1 | 1s9j_v | 9 | 3.811 | 0.4234 | 0.82871 | 7 | 2 | 0 | 0 | 0 | 0 | 0 | 0 | Dual spec | TRANSFER | AMP2K1_HUM | Catalyzes | Defects iColon tum | |
| SETD7 | 1mt6_v | 7 | 2.961 | 0.423 | 0.571635 | 2 | 2 | 2 | 0 | 0 | 1 | 0 | 0 | Histone-1 | TRANSFER | SETD7_HUM | Histone | #NONE | |
| HCK | 1qcf_v | 7 | 2.945 | 0.4207 | -0.0957678 | 3 | 3 | 1 | 0 | 0 | 0 | 0 | 0 | Tyrosine | -TYROSINE | HCK_HUMAN | May serve | NONE | |
| S100A9 | 1irj_v | 7 | 2.939 | 0.4198 | 0.252421 | 4 | 0 | 3 | 0 | 0 | 0 | 0 | 0 | Protein | SMETAL | BINS10A9_HUM | Expressed | NONE | |
| TEK | 2oo8_v | 8 | 3.35 | 0.4188 | 0.714277 | 5 | 1 | 1 | 1 | 0 | 0 | 0 | 0 | Angiopoie | TRANSFER | TE2_HUM | This prot | Defects i | |
| ACPP | 1nd5_v | 6 | 2.509 | 0.4181 | -1.06311 | 2 | 2 | 0 | 0 | 1 | 1 | 0 | 0 | Prostatic | HYDROLASE | PPAP_HUMAN | None | Prostate | |
| HSP90AB1 | 1uyu_v | 8 | 3.341 | 0.4176 | 1.40731 | 5 | 1 | 1 | 0 | 0 | 0 | 0 | 1 | Heat shoc | None | P08238 | Posttrans | NONE | |
| KAT2B | 1cm0_v | 9 | 3.723 | 0.4137 | 2.8146 | 1 | 6 | 2 | 0 | 0 | 0 | 0 | 0 | Histone | aNONE | Q92831 | Involved | NONE | |
| TK1 | 1w4r_v | 6 | 2.482 | 0.4136 | -0.113883 | 0 | 2 | 2 | 0 | 0 | 2 | 0 | 0 | Thymidine | None | P04183 | Nucleotid | NONE | |
| ERBB4 | 3bbt_v | 6 | 2.476 | 0.4126 | -0.925959 | 5 | 1 | 0 | 0 | 0 | 0 | 0 | 0 | Receptor | TRANSFER | ERBB4_HUM | Specific | NONE | |
| CTSF | 1md6_v | 7 | 2.88 | 0.4115 | 0.384493 | 3 | 2 | 2 | 0 | 0 | 0 | 0 | 0 | Cathepsin | None | Q9UBX1 | Involved | NONE | |
| SDS | 1p5j_v | 7 | 2.877 | 0.411 | -0.188348 | 2 | 4 | 0 | 0 | 1 | 0 | 0 | 0 | L-serine | None | P20132 | Amino aci | NONE | |
| PSAP | 1n69_v | 7 | 2.876 | 0.4108 | -0.0666955 | 6 | 0 | 0 | 0 | 0 | 1 | 0 | 0 | Proactiva | None | P07602 | Involved | NONE | |
| PAPPS1 | 1x6v_v | 7 | 2.874 | 0.4106 | -0.194749 | 0 | 5 | 1 | 0 | 0 | 1 | 0 | 0 | Bifunctio | None | Q43252 | Involved | NONE | |
| #N/A | 1svh_v | 7 | 2.871 | 0.4101 | 0.322527 | 4 | 1 | 1 | 1 | 0 | 0 | 0 | 0 | cAMP-depe | None | P00517 | Involved | NONE | |
| NQO2 | 1sgo_0_v | 5 | 2.045 | 0.409 | -1.39821 | 2 | 2 | 0 | 0 | 0 | 0 | 0 | 0 | Ribosyldi | OXIDOREDU | NQO2_HUM | The enzym | NONE | |
| CTSS | 1nqc_v | 7 | 2.841 | 0.4059 | 0.119659 | 3 | 2 | 2 | 0 | 0 | 0 | 0 | 0 | Cathepsin | HYDROLASE | CATS_HUM | Thiol prc | Psoriasis | |
| MMP16 | 1rm8_v | 8 | 3.236 | 0.4045 | 0.887687 | 1 | 2 | 4 | 0 | 0 | 0 | 0 | 1 | Matrix | meHYDROLASE | MMP16_HUM | Endopepti | NONE | |
| MAN1B1 | 1x9d_v | 7 | 2.829 | 0.4041 | 0.183273 | 0 | 5 | 2 | 0 | 0 | 0 | 0 | 0 | Endoplasm | None | Q9U1M7 | Involved | NONE | |
| NT5M | 1q91_v | 9 | 3.634 | 0.4038 | 1.09764 | 1 | 6 | 1 | 0 | 1 | 0 | 1 | 0 | 5 (3)-deox | None | Q9NPB1 | Involved | NONE | |
| ACADM | legc_v | 7 | 2.823 | 0.4033 | 0.479859 | 3 | 2 | 2 | 0 | 0 | 0 | 0 | 0 | Medium-chELECTRON | ACADM_HUM | This enzym | Defects i | | |
| TGM3 | 1ln9n_v | 8 | 3.199 | 0.3998 | 0.0664704 | 3 | 4 | 1 | 0 | 0 | 0 | 0 | 0 | Protein-g | TRANSFER | TGM3_HUM | Catalyzes | NONE | |
| DTYMK | 1mmx_v | 7 | 2.785 | 0.3979 | -0.28059 | 1 | 4 | 1 | 0 | 0 | 1 | 0 | 0 | Thymidyla | TRANSFER | KTHY_HUM | Catalyzes | NONE | |
| GSTM1 | 1xwk_v | 9 | 3.58 | 0.3978 | 1.96215 | 2 | 2 | 3 | 1 | 1 | 1 | 0 | 0 | Glutathio | TRANSFER | GSTM1_HUM | Conjugati | NONE | |
| CSK | 3dt7_v | 6 | 2.387 | 0.3978 | -0.752325 | 3 | 1 | 1 | 1 | 0 | 0 | 0 | 0 | Tyrosine | -TRANSFER | CSK_HUMAN | Specific | NONE | |
| AGXT | 1h0c_v | 7 | 2.77 | 0.3958 | -0.350289 | 2 | 2 | 1 | 0 | 1 | 0 | 1 | 1 | Serine- | fTRANSFER | SPYA_HUMAN | Defects iHyperoxal | | |
| SHMT1 | 1bj4_v | 6 | 2.368 | 0.3946 | -0.676854 | 2 | 4 | 0 | 0 | 0 | 0 | 0 | 0 | Serine | hyTRANSFER | AGLYC_HUM | Interconv | NONE | |
| IMPDH1 | 1jcn_v | 7 | 2.753 | 0.3933 | -0.363561 | 0 | 5 | 1 | 0 | 0 | 1 | 0 | 0 | Inosine-50 | XDOREDU | IMDH1_HUM | Rate limi | Defects iNONE | |
| HNMT | 1jqe_v | 8 | 3.136 | 0.392 | 0.00536757 | 7 | 0 | 0 | 1 | 0 | 0 | 0 | 0 | Histamine | None | P50135 | Inactivat | NONE | |
| PPCDC | 1qzu_v | 7 | 2.73 | 0.39 | 0.134535 | 1 | 3 | 2 | 0 | 0 | 1 | 0 | 0 | Phosphopan | eNONE | Q96CD2 | Involved | NONE | |
| HAGH | 1qh5_v | 7 | 2.692 | 0.3846 | -0.331195 | 0 | 4 | 1 | 0 | 2 | 0 | 0 | 0 | Hydroxyac | None | Q16775 | Involved | NONE | |
| RXR8 | 1lh9u_v | 10 | 3.781 | 0.3781 | 0.716892 | 8 | 1 | 0 | 0 | 0 | 1 | 0 | 0 | 0 | Retinoic | NUCLEAR | RXR8B_HUMAN | Nuclear | NONE |
| LGALS2 | 1hlc_v | 6 | 2.267 | 0.3778 | -1.13758 | 0 | 3 | 3 | 0 | 0 | 0 | 0 | 0 | Galectin | -LECTIN | LEG2_HUM | This prot | NONE | |
| CMA1 | 1t31_v | 8 | 3.021 | 0.3776 | 0.267583 | 3 | 4 | 0 | 0 | 0 | 1 | 0 | 0 | Chymase | HYDROLASE | CMA1_HUM | Major sec | NONE | |
| MME | 1r1l_v | 8 | 2.997 | 0.3746 | 0.0692056 | 3 | 2 | 1 | 1 | 1 | 1 | 0 | 0 | Nepriyls | HYDROLASE | NEF_HUMAN | Thermolys | Important Hypertens | |
| CCBL1 | 1w7n_v | 6 | 2.242 | 0.3736 | -1.64553 | 1 | 2 | 1 | 1 | 1 | 1 | 0 | 0 | Kynurenin | TRANSFER | AKT1_HUM | Catalyzes | NONE | |
| CD209 | 1s14_v | 8 | 2.956 | 0.3695 | 0.898312 | 0 | 5 | 3 | 0 | 0 | 0 | 0 | 0 | CD209 | ant SUGAR | BINC209_HUM | Pathogen | NONE | |
| AKT2 | 2uw9_v | 6 | 2.214 | 0.369 | -0.979077 | 3 | 1 | 1 | 1 | 0 | 0 | 0 | 0 | RAC-beta | TRANSFER | AKT2_HUM | General | fAlteratio | |
| PPP1CC | 1jk7_v | 9 | 3.314 | 0.3682 | 0.736053 | 6 | 1 | 1 | 0 | 1 | 0 | 0 | 0 | Serine/th | None | P36873 | Signal tr | NONE | |
| MMP1 | 1hfc_v | 8 | 2.943 | 0.3678 | 0.176404 | 1 | 3 | 4 | 0 | 0 | 0 | 0 | 0 | Interstiti | None | P03956 | Cleaved | cNONE | |
| F11 | 1zpb_v | 8 | 2.934 | 0.3668 | 0.346602 | 3 | 3 | 1 | 1 | 0 | 0 | 0 | 0 | Coagulati | HYDROLASE | FA11_HUM | Factor X | Defects iThrombosi | |
| BCAT2 | 1ekv_v | 8 | 2.932 | 0.3665 | -0.0374593 | 2 | 3 | 2 | 0 | 0 | 1 | 0 | 0 | Branched- | NONE | 015382 | Amino aci | NONE | |
| LGALS3 | 1kjl_v | 7 | 2.558 | 0.3654 | -0.403225 | 0 | 3 | 4 | 0 | 0 | 0 | 0 | 0 | Galectin | -SUGAR | BINLEG3_HUM | Galactose | Colorecta | |
| MMP9 | 1gkc_v | 8 | 2.9 | 0.3626 | -0.134847 | 2 | 4 | 2 | 0 | 0 | 0 | 0 | 0 | Matrix | meHYDROLASE | MMP9_HUM | play -Leu | bondDefects i | |
| GNPDA1 | 1ne7_v | 7 | 2.53 | 0.3614 | -0.719854 | 0 | 4 | 3 | 0 | 0 | 0 | 0 | 0 | Glucosami | None | P46926 | Carbohydr | NONE | |
| UCK2 | 1uj2_v | 8 | 2.877 | 0.3597 | -0.234299 | 1 | 3 | 3 | 0 | 1 | 0 | 0 | 0 | Uridine-c | None | Q9BZK2 | Nucleotid | NONE | |
| CLEC4M | 1k9j_v | 8 | 2.857 | 0.3572 | 0.779822 | 0 | 5 | 3 | 0 | 0 | 0 | 0 | 0 | C-type le | None | Q9H2X3 | Probable | NONE | |
| GP1BA | 1p9a_v | 8 | 2.806 | 0.3507 | 0.327505 | 0 | 5 | 3 | 0 | 0 | 0 | 0 | 0 | Platelet | None | P07359 | Involved | NONE | |
| GSTO1 | 1eev_v | 8 | 2.794 | 0.3492 | -0.131799 | 0 | 4 | 2 | 1 | 1 | 1 | 0 | 0 | Glutathio | TRANSFER | GSTO1_HUM | Exhibits | NONE | |
| STAT1 | 1yv1_v | 8 | 2.79 | 0.3488 | -0.108378 | 1 | 4 | 2 | 1 | 0 | 0 | 0 | 0 | Signal tr | SIGNALING | STAT1_HUN | Signal tr | Defects iRespirato | |
| CDC42 | 1a4r_v | 8 | 2.786 | 0.3483 | -0.461036 | 0 | 5 | 2 | 0 | 0 | 1 | 0 | 0 | Cell divi | HYDROLASE | CDC42_HUM | Plasma | meNONE | |
| CTS8 | 1gmy_v | 9 | 3.134 | 0.3482 | 0.396921 | 4 | 2 | 3 | 0 | 0 | 0 | 0 | 0 | Cathepsin | COMPLEX | (CATB_HUM) | Thiol | Myocardia | |
| PTCP | 1ln3_v | 11 | 3.806 | 0.346 | 1.13577 | 9 | 0 | 0 | 1 | 1 | 1 | 0 | 0 | Phosphati | None | Q9UKL6 | Involved | NONE | |
| GPI | 1nuh_v | 8 | 2.753 | 0.3442 | -0.645543 | 1 | 5 | 1 | 0 | 0 | 1 | 0 | 0 | Glucose-6 | ISOMERASE | GP1_HUMAN | Neurotrop | Defects iNONE | |
| #N/A | 2qcf_v | 6 | 2.065 | 0.3442 | -1.58237 | 0 | 3 | 2 | 0 | 1 | 1 | 0 | 0 | Uridine 5 | LYASE | PYR5_HUMAN | Defects | iSquamous | |
| OTC | 1oth_v | 8 | 2.741 | 0.3426 | -0.527379 | 1 | 4 | 1 | 1 | 1 | 1 | 0 | 0 | Ornithine | None | P00480 | Amino aci | NONE | |
| TPSB2 | 2fs8_v | 8 | 2.737 | 0.3421 | 0.0670373 | 2 | 2 | 2 | 1 | 0 | 1 | 0 | 0 | Tryptase | HYDROLASE | TRYVB2_HUM | Tryptase | NONE | |
| ARF4 | 1z6x_v | 8 | 2.735 | 0.3419 | -0.438133 | 0 | 4 | 2 | 0 | 2 | 0 | 0 | 0 | ADP- | | | | | |

ase;Musculoskeletal disease;Periodontal disease;Bone injury;Osteoporosis
fibroids;Menstruation disorder;Infertility;Acne;Breast tumor;Female contraception;Neoplasm;Planned abortion;Dysmenorrhea;Endometriosis;Hormone deficiency;Unidentif

disorder;Cancer;Breast tumor

city, drug-induced;Toxicity;Alzheimers disease
tumor
Cancer;Liver disease;Cognitive disorder;Neurodegenerative disease;Cerebrovascular ischemia
NONE
tional supplementation, also for treating dietary shortage or imbalance

Brain tumor;Inflammation;Myocardial infarction;Fibrosis;Brain ischemia;Psoriasis;Rheumatoid arthritis;Retinitis pigmentosa;Atherosclerosis;Genitourinary disease;B
y;Diabetic neuropathy;Diabetic foot ulcer;Diabetic complication;Diabetic nephropathy;Retinopathy;Ocular disease;Central nervous system disease;Diabetic retinopath
ment of transplant rejection, rheumatoid arthritis, severe psoriasis

obstructive pulmonary disease;Asthma

tional supplementation, also for treating dietary shortage or imbalance

Hodgkins disease;Acute leukemia;Breast tumor;Head and neck tumor;Hepatocellular carcinoma;Ovary tumor;Prostate tumor;Solid tumor;Non-Hodgkin lymphoma;Cancer;Non-s
ervous system disease;Inflammation;Pulmonary fibrosis;Ischemia;Cerebrovascular ischemia;Neurodegenerative disease

;Asthma;Cancer;Neurodegenerative disease;Inflammation;Liver disease;Acute myelogenous leukemia;Diabetes mellitus;Pulmonary fibrosis;Ischemia
s disease;Cocaine addiction;Major depressive disorder;Epilepsy;Neurodegenerative disease;Attention deficit hyperactivity disorder;Head injury;Parkinsons disease;D
mphoma;Autoimmune disease;HIV infection;Dermatitis;Chronic lymphocytic leukemia;Acquired Immune Deficiency Syndrome;Inflammation;Multiple sclerosis;Solid tumor;Ps
e sexual desire disorder;Hepatitis B virus infection;Cardiac failure;Andrology;Hirsutism;Female sexual dysfunction;Psoriasis;Reperfusion injury;Plasmodium infecti
lerosis;Cerebrovascular disease;Cancer;Melanoma;Multiple myeloma;Hypercholesterolemia;Atherosclerosis;Breast tumor;Hyperlipidemia;Solid tumor;Cardiovascular disea
onuria;Sickle cell anemia;Vascular disease;Genetic disorder;Diabetic nephropathy;Hypertension;Ataxia;Peripheral arterial occlusive disease;Coronary artery disease

lence suggests that NADH might be useful in treating Parkinsons disease, chronic fatigue syndrome, Alzheimers disease and cardiovascular disease.

is;Glomerulonephritis;Wound healing;Renal disease;Scleroderma;Glaucoma;Diabetic complication;Scar tissue;Inflammation;Urinary tract disease;Pulmonary fibrosis;Diab
disorder;Hypertension;Cardiovascular disease;Urinary tract disease

demia;Hypercholesterolemia;Atherosclerosis
emia;Cancer;Osteoporosis;Cerebrovascular ischemia;Bone metastases;Metastasis;Solid tumor
in dependent diabetes

cell carcinoma;Arteriosclerosis;Melanoma;Central nervous system tumor;Diabetes mellitus;Esophagus tumor;Ischemic heart disease;Brain tumor;Nasopharyngeal carcinom
solid tumor
Renal disease;Cardiac failure;Atherosclerosis;HIV infection;Hypertension;Cardiovascular disease;Ocular hypertension;Renal failure
artery disease;Hypertension;Congestive heart failure;Angina;Myocardial infarction;Peripheral vascular disease;Atherosclerosis;Ischemic heart disease

tional supplementation, also for treating dietary shortage or imbalance
l, known for its antioxidant activities, is protective against cardiovascular disease and some forms of cancer and has also demonstrated immune-enhancing effects.
;Inflammation;Rheumatoid arthritis;Cardiac failure;Multiple sclerosis;Metastasis;Immune disorder
be used for lowering elevated triglycerides in those who are hyperglyceridemic. In addition, EPA may play a therapeutic role in patients with cystic fibrosis by re

in dependent diabetes;Rheumatoid arthritis;Diabetes mellitus;Obesity;Atherosclerosis;Metabolic disorder;Periodontitis
rebrovascular ischemia;Conjunctivitis;Inflammation;Multiple sclerosis;Fibrosis;Unidentified indication;Psoriasis;Heart arrhythmia;Rheumatoid arthritis;Atheroscler
dependent diabetes;Glomerulonephritis;Cancer;Rheumatoid arthritis;Sepsis;Autoimmune disease;Atherosclerosis;Infection;Inflammation;Myocardial infarction;Immune dis
sis disorder;Melanoma;Cancer;Non-small-cell lung cancer;Renal cell carcinoma;Colorectal tumor;Ocular disease;Age related macular degeneration;Solid tumor
lung cancer;Solid tumor;Breast tumor;Head and neck tumor;Brain tumor;Neoplasm;Hepatocellular carcinoma;Ovary tumor;Prostate tumor;Pancreas tumor;Urinary tract t
Cancer

myelocytic leukemia
complication

sclerosis;Asthma;Cancer;Rheumatoid arthritis;Autoimmune disease;Arthritis;Bacterial infection;Bone disease;Osteoarthritis;Bone metastases;Immune disorder;Osteopor

ment of diabetes (type I and II)

tional supplementation, also for treating dietary shortage or imbalance

: term treatment of acutely decompensated severe chronic heart failure (CHF). Also being investigated for use/treatment in heart disease.

abolism

estigated for the treatment of Herpes labialis infections (cold sores).

, known for its antioxidant activities, is protective against cardiovascular disease and some forms of cancer and has also demonstrated immune-enhancing effects. tional supplementation and for treating dietary shortage or imbalance.

d arthritis;Diabetes mellitus;Non-insulin dependent diabetes;Autoimmune disease;Immune disorder

;Cancer;Rheumatoid arthritis;Cardiac failure;Atherosclerosis;Delayed hypersensitivity;Transplant rejection;Inflammation;Multiple sclerosis;Autoimmune disease

ne disease;Reperfusion injury;Age related macular degeneration
e tissue disease;Viral infection;Inflammation;HIV infection

hy;Vascular disease;Diabetic macular edema;Inflammation;Cancer;Gastrointestinal disease;Age related macular degeneration

anthracis infection

i thrombosis;Thrombocytopenia;Cerebrovascular disease;Wound healing;Ebola virus infection;Bone injury;Cerebrovascular ischemia;Restenosis;Thrombosis;Diabetic foot ion;Cancer;Acute myelogenous leukemia;Myocardial infarction;Non-small-cell lung cancer;Solid tumor ure;Raynauds disease;Cardiac failure;Thrombosis;Diabetic complication;Prostate hyperplasia;Erectile dysfunction;Urinary tract disease;Female sexual dysfunction;Va

sis disorder;Ulcer;Cancer;Lung embolism;Restenosis;Metastasis;Skin ulcer;Cardiovascular disease
obstructive pulmonary disease
s disease;Renal disease;Lipid metabolism disorder;Cardiac failure;Hyperlipidemia;Intermittent claudication;Myocardial infarction;Coronary artery disease;Glioma;Hy

lence suggests that NADH might be useful in treating Parkinsons disease, chronic fatigue syndrome, Alzheimers disease and cardiovascular disease.
ion;Cancer

;Genital tract inflammation;Otitis media;Wound healing;Atopic dermatitis;Dermatitis;Cystitis;Inflammatory bowel disease;Cystic fibrosis;Alpha-1 antitrypsin defici se;Rheumatoid arthritis;Cancer

;Transplant rejection;Rheumatoid arthritis;Respiratory tract inflammation;Autoimmune disease

ildrich syndrome

abolism disorder;Atherosclerosis
ment of mucositis (mouth sores)

X;Lipid metabolism disorder;Non-insulin dependent diabetes;Atherosclerosis;Hypertriglyceridemia ;Hyperlipidemia;Cardiovascular disease;Congestive heart failure;Dia tional supplementation, also for treating dietary shortage or imbalance.
rosis;Non-insulin dependent diabetes;Atherosclerosis;Hypertriglyceridemia ;Hyperlipidemia;Liver disease;Primary biliary cirrhosis;Non-alcoholic steatohepatitis lence suggests that NADH might be useful in treating Parkinsons disease, chronic fatigue syndrome, Alzheimers disease and cardiovascular disease.

replacement or supplemental therapy in patients with hypothyroidism of any etiology, except transient hypothyroidism during the recovery phase of subacute thyroidit lupus erythematosus;Asthma;Rheumatoid arthritis;Allergy;Allergic rhinitis;Inflammation;Thrombocytopenic purpura;Acute myelogenous leukemia;Lymphoma

tional supplementation, also for treating dietary shortage or imbalance
in tumor;Leukemia;Breast tumor;Lung tumor;Head and neck tumor;Uterine cervix tumor;Ovary tumor;Prostate tumor;Prostate hyperplasia

is claimed to act as an effective antidepressant, however results are mixed. Tyrosine has also been claimed to reduce stress and combat narcolepsy and chronic fat

drug-induced;Mucositis;Cancer;Glioma;Metastasis;Renal cell carcinoma;Head and neck tumor;Brain tumor;Pancreas tumor;Solid tumor;Stomach tumor

;thrombosis;Blood clotting disorder;Cancer;Ebola virus infection;Thrombosis;Lung embolism;Unstable angina;Angina;Myocardial infarction;Thromboembolism;Cardiovascu treatment of ariboflavinosis (vitamin B2 deficiency).
;treatment of hypercholesterolemia.

Cancer;Solid tumor

s disease;Immune deficiency;Breast tumor;Major depressive disorder;Glaucoma;Metabolic disorder;Psychotic disorder;HIV infection;Cocaine addiction;Inflammation;Pla me disease;Melanoma;Hyperthyroidism;Aging;Skin tumor;Acne;Breast tumor;Neurodegenerative disease;Acute myelogenous leukemia;Hepatocellular carcinoma;Prostate tumo ill anemia;Inflammation;Transplant rejection;Myocardial infarction;Restenosis;Thrombosis
l tumor;Non-Hodgkin lymphoma;Neoplasm;Cancer;Trypanosomiasis;Pneumocystis carinii infection;Bacterial infection

ical neoplasm;Multiple myeloma

al treatment of cutaneous lesions in patients with AIDS-related Kaposi's sarcoma.

n adjunct to the standard therapy of inhaled steroids with inhaled long- and/or short-acting beta-agonists.
tional supplementation, also for treating dietary shortage or imbalance
;Cancer;Rheumatoid arthritis;Pulmonary hypertension;Myeloid leukemia;Inflammation;Prostate tumor;Pancreas tumor;Myeloproliferative disorder;Solid tumor;Myelofibro ositis;Migraine;Neuropathic pain;Irritable bowel syndrome;Reperfusion injury;Inflammatory bowel disease;Inflammation;Chronic obstructive pulmonary disease;Gasti tritis;Chronic bronchitis;Lung injury;Cerebrovascular ischemia;Emphysema;Respiratory disease;Cystic fibrosis;Chronic obstructive pulmonary disease;Myocard

tional supplementation, also for treating dietary shortage or imbalance

ion

;Acne;Cancer

cell carcinoma;Psoriasis;Colon tumor;Cancer;Metabolic disorder;Leukemia;Breast tumor;Skin infection;Neutropenia;Carcinoma;Dermatological disease;Ocular disease;Ge
t rejection;Immune disorder

NONE

or;Melanoma;Cancer;Rheumatoid arthritis;Non-small-cell lung cancer;Neurodegenerative disease;Breast tumor;Colorectal tumor;Transplant rejection;Inflammation;Pancre

hy;Ovary tumor;Cancer;Renal cell carcinoma;Breast tumor;Solid tumor
tumor

tional supplementation and for treating dietary shortage or imbalance.

;Arteriosclerosis;Pain;Rheumatoid arthritis;Autoimmune disease;Allergy;Inflammation;Asthma;Multiple sclerosis;Immune disorder

uria

ment of acute malarial attacks in non-immune subjects.

tional supplementation, also for treating dietary shortage or imbalance

opic dermatitis;Atherosclerosis;Thrombosis;Bacterial infection;Fungal infection;Inflammation;Congestive heart failure;Chronic obstructive pulmonary disease;Viral
infection;Neuropathic pain;Renovascular hypertension;Congestive heart failure

s
tional supplementation and for treating dietary shortage or imbalance.
l tumor;Chronic lymphocytic leukemia;Cancer;Prostate tumor;Pancreas tumor;Metastasis;Multiple myeloma
NONE

ry disease;Asthma

l infarction;Arthritis;Inflammation;Cancer

cell carcinoma;Biliary cancer;Cancer;Non-small-cell lung cancer;Gastrointestinal tumor;Renal cell carcinoma;Breast tumor;Lung tumor;Head and neck tumor;Colorectal
nutritional supplementation, also for treating dietary shortage or imbalance. It has been claimed that ornithine improves athletic performance, has anabolic effec

Breast tumor;Cancer

replacement or supplemental therapy in patients with hypothyroidism of any etiology, except transient hypothyroidism during the recovery phase of subacute thyroidit

tional supplementation, also for treating dietary shortage or imbalance

Metabolic disorder;Neurodegenerative disease;Cancer;Cardiovascular disease

ovirus infection;Metastasis;Cancer;Solid tumor
ing injury;Atopic dermatitis;Cerebrovascular ischemia;Shock;Ischemic heart disease;Dermatitis;Respiratory disease;Inflammatory bowel disease;Inflammation;Chronic o
neck tumor;Carcinoma;Melanoma;Cancer;Sarcoma;Non-small-cell lung cancer;Breast tumor
tumor;Ovary tumor;Cancer;Mastocytosis;Gastrointestinal tumor

;Septic shock;Alzheimers disease;Rheumatoid arthritis;Sepsis;Neurological disease;Osteoarthritis;Inflammation;Diabetes mellitus;Metastasis;Cardiovascular disease

ied indication;Gynecological disorder;Musculoskeletal disease;Estrogen deficiency;Cancer;Contraception;Amenorrhea;Female infertility;Carcinoma;Menopause;Premenstr

breast disease; Atrophy; Crohn's disease; Testosterone deficiency; Uterine fibroids; Fibrocystic breast disease; Arthralgia; Prostatitis; Prostate hyperplasia; Inflammatory bowel disease; Diabetes mellitus; Diabetic cataract; Cataract

small-cell lung cancer;Renal cell carcinoma;Colorectal tumor

Dementia; Nicotine dependence; Cerebrovascular ischemia

oriasis;Non-Hodgkin lymphoma;Viral infection;Cancer;Rheumatoid arthritis;Leukemia;B-cell acute lymphoblastic leukemia;Cutaneous T-cell lymphoma;Ocular disease;Trauma;Atherosclerosis;Prostate hyperplasia;Fatigue;Male sexual dysfunction;Autoimmune disease;Testosterone deficiency;Uterine fibroids;Metabolic disorder;Myalgia;Obesity;Prostate tumor;Alzheimers disease;Dementia;Osteoporosis

etic nephropathy; Fibrosis; Solid tumor; Angiogenesis disorder; Glioma; Cancer; Atherosclerosis; Glomerular disease; Lung tumor; Nephritis; Immune disorder; Bone marrow trans

a:Liver tumor;Pancreas tumor;Lung tumor;Psoriasis;Bladder tumor;Esophageal disease;Mesothelioma;Atherosclerosis;Multiple myeloma;Bacterial infection;Uterine cervix

It may be of limited benefit in some with asthma and rheumatoid arthritis. It may be helpful in some neurological diseases including Alzheimers, some eye disorders producing disease severity and may play a similar role in type 2 diabetics in slowing the progression of diabetic nephropathy.

osis;Corneal disease;Cardiovascular disease;Ulcerative colitis;Alzheimers disease;Infertility;Syndrome X;Lipid metabolism disorder;Non-insulin dependent diabetes;order

umor:Stomach tumor:Retinopathy:Bladder tumor:Colon tumor:Cancer:Mesothelioma:Non-small-cell lung cancer:Cholangiocarcinoma:Gastrointestinal tumor:Liver tumor:Lung

It may be of limited benefit in some with asthma and rheumatoid arthritis. It may be helpful in some neurological diseases including Alzheimers, some eye disorders:

ulcer;Ischemic heart disease;Marburg virus infection;Myocardial disease;Embolism and thrombosis;Myocardial infarction;Cartilage disease;Thromboembolism;Coronary artery disease;Viral infection;Cancer;Atherosclerosis;Pulmonary disease;Metabolic disorder;Hypertension;Cardiovascular disease;Pulmonary hypertension;Angina;Male :
percholesterolemia ;Atherosclerosis ;Peripheral arterial occlusive disease;Cardiovascular disease;Congestive heart failure;Angina;Hypertension;Heart disease;Cerebro-

ency;Dermatological disease

abetes mellitus;Obesity;Inflammation

is.

igue, however these claims have been refuted by some studies.

lar disease

nned abortion;Diabetes mellitus;Weight gain;Endometriosis;Obesity;Insulin dependent diabetes;Cushings disease;Glioma;Endometroid carcinoma;Neurodegenerative disease;Pancreas tumor;Alzheimers disease;Multiple sclerosis;Psoriasis;Colon tumor;Cancer;Renal disease;Leukemia;Hyperparathyroidism;Sarcoidosis;Carcinoma;Insulin dependence;

sis;Hematological neoplasm;Cardiovascular disease
c motility disorder;Uveitis;Insulin dependent diabetes;Multiple sclerosis;Rheumatoid arthritis;Sepsis;Osteoarthritis;Transplant rejection;Gingivitis;Allergic rhinitis;Arterial infarction;Lung inflammation;Pulmonary fibrosis;Psoriasis;Pancreatitis;Bronchitis;Rheumatoid arthritis;Arthritis;Pulmonary disease;Lung tumor;Respiratory distress;

nital system disease;Lymphoma;Immune disorder

eas tumor;Solid tumor

infection;HIV infection;Cardiovascular disease

tumor;Adenocarcinoma;Prostate tumor;Pancreas tumor;Hepatocellular carcinoma;Stomach tumor
ts, has wound-healing effects, and is immuno-enhancing.

is.

bstructive pulmonary disease;Myocardial infarction;Injury;Psoriasis;Reperfusion injury;Sepsis;Skin burns;Respiratory distress syndrome;Transplant rejection;Hypertension

al syndrome;Anesthesia;Age related macular degeneration;Osteoporosis

bowel disease;Solid tumor;Gynecological disorder;Vascular disease;Contraception;Cancer;Sepsis;Hypercholesterolemia;Neurological disease;Ataxia;Cognitive disorder;(

nsplant rejection;Immune disorder
sity;Gynecological disorder;Contraception;Cancer;Hypercholesterolemia;Skin burns;Seborrhea;Growth disorder;Bone tumor;Muscular dystrophy;Erectile dysfunction;Syste

splantation

x tumor;Nasopharynx tumor;Carcinoma;Restenosis;Stomach tumor;Thyroid tumor;Small-cell lung cancer;Renal disease;Nervous system tumor;Hepatobiliary system tumor;Bre

s including cataracts, and diabetes and premenstrual syndrome. It may also help protect skin from ultraviolet irradiation although claims that it reverses skin agi

Metabolic disorder;Diabetic complication;Hyperlipidemia;Lacrimal gland disease;Myocardial infarction;Obesity;Coronary artery disease;Insulin dependent diabetes;Vas

tumor;Colorectal tumor;Uterine cervix tumor;Carcinoma;Biliary tumor;Gallbladder disease;Keratosis

s including cataracts, and diabetes and premenstrual syndrome. It may also help protect skin from ultraviolet irradiation although claims that it reverses skin agi

rtery disease;Hemophilia;Blood clotting disorder;Bleeding;Atrial fibrillation;Skin burns;Disseminated intravascular coagulation;Connective tissue disease;Lung embc
sexual dysfunction

vascular ischemia;Osteoporosis

se;Hypertension;Hepatitis C virus infection;Carcinoma;Non-insulin dependent diabetes;Psychiatric disorder
dent diabetes;Transplant rejection;Osteoporosis

itis

ress syndrome;Transplant rejection;Pulmonary hypertension;Pneumonia;Inflammation

Osteoporosis;Gynecomastia;Major depressive disorder;Acne;Female contraception;Dysmenorrhea;Neoplasm;Central nervous system disease;Multiple sclerosis;Alcoholism;La

emic lupus erythematosus;Asthma;Wound healing;Cerebrovascular ischemia;Acne;Hypogonadism;HIV infection;Neoplasm;Dermatological disease;Prostate disease;Multiple sc

east tumor;Head and neck tumor;Ovary tumor;Prostate tumor;Renal tumor;Solid tumor;Colon tumor;Glioma;Cancer;Non-small-cell lung cancer;Metastasis;Renal cell carcin

ing, enhances male fertility and exercise performance are poorly supported. It may help relieve some muscle cramps.

scular disease;Cancer;Cognitive disorder;Hypertension;Diabetes mellitus;Osteoporosis;Dermatological disease

ing, enhances male fertility and exercise performance are poorly supported. It may help relieve some muscle cramps.

olism;Cardiovascular disease;Sepsis;Angina;Infarction

lacrimal gland disease;Carcinoma;Cardiovascular disease;Menopause;Unidentified indication;Alzheimers disease;Breast tumor;Scar tissue;Hyperlipidemia;Postmenopausal

lesterol;Lacrimal gland disease;Alopecia;Mycobacterium tuberculosis infection;Ovary cyst;Carcinoma;Cardiovascular disease;Cachexia;Muscle wasting disease;Cervical

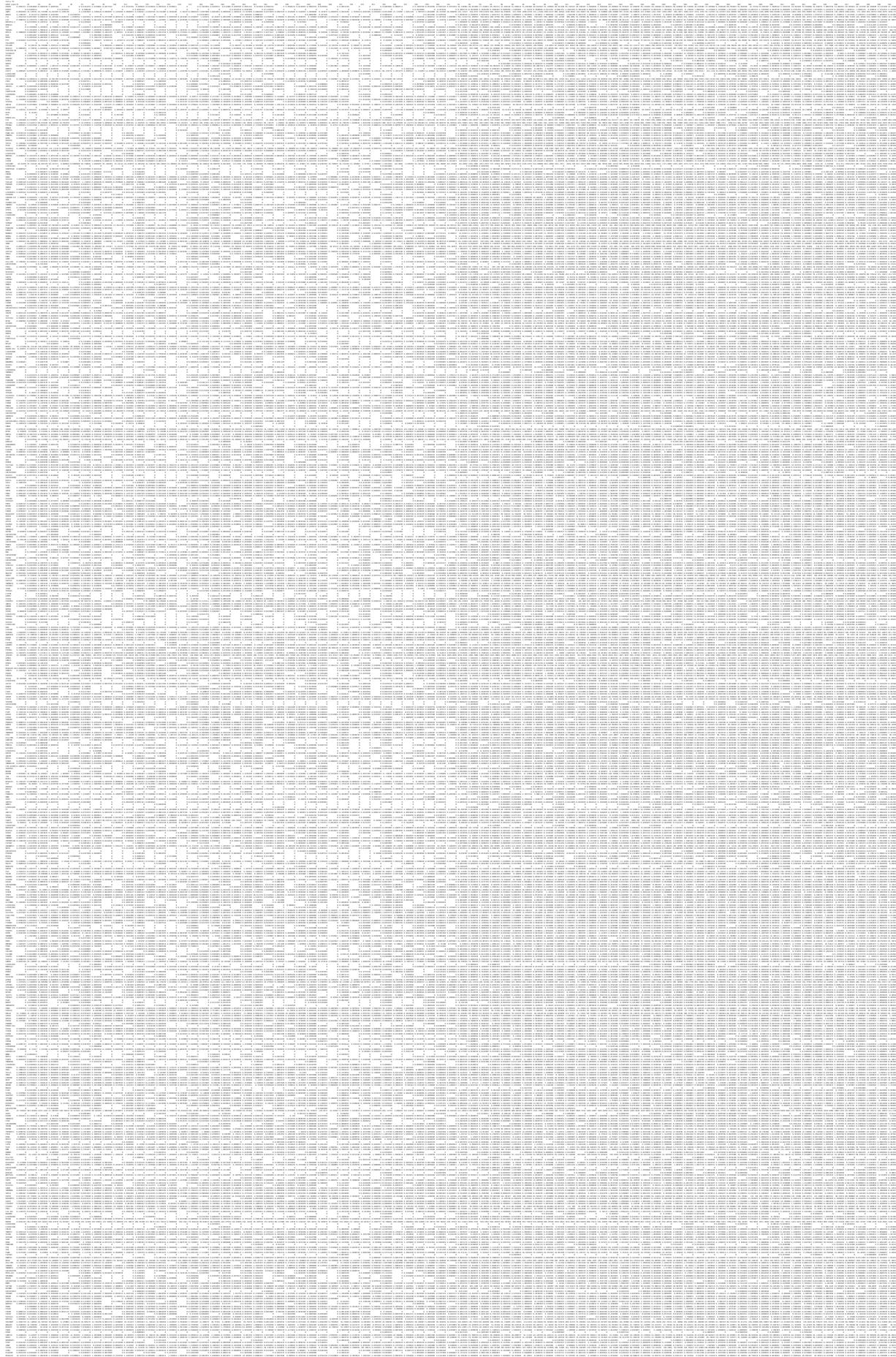
oma;Glioblastoma;Colorectal tumor;Neoplasm;Sarcoma;Hypercholesterolemia

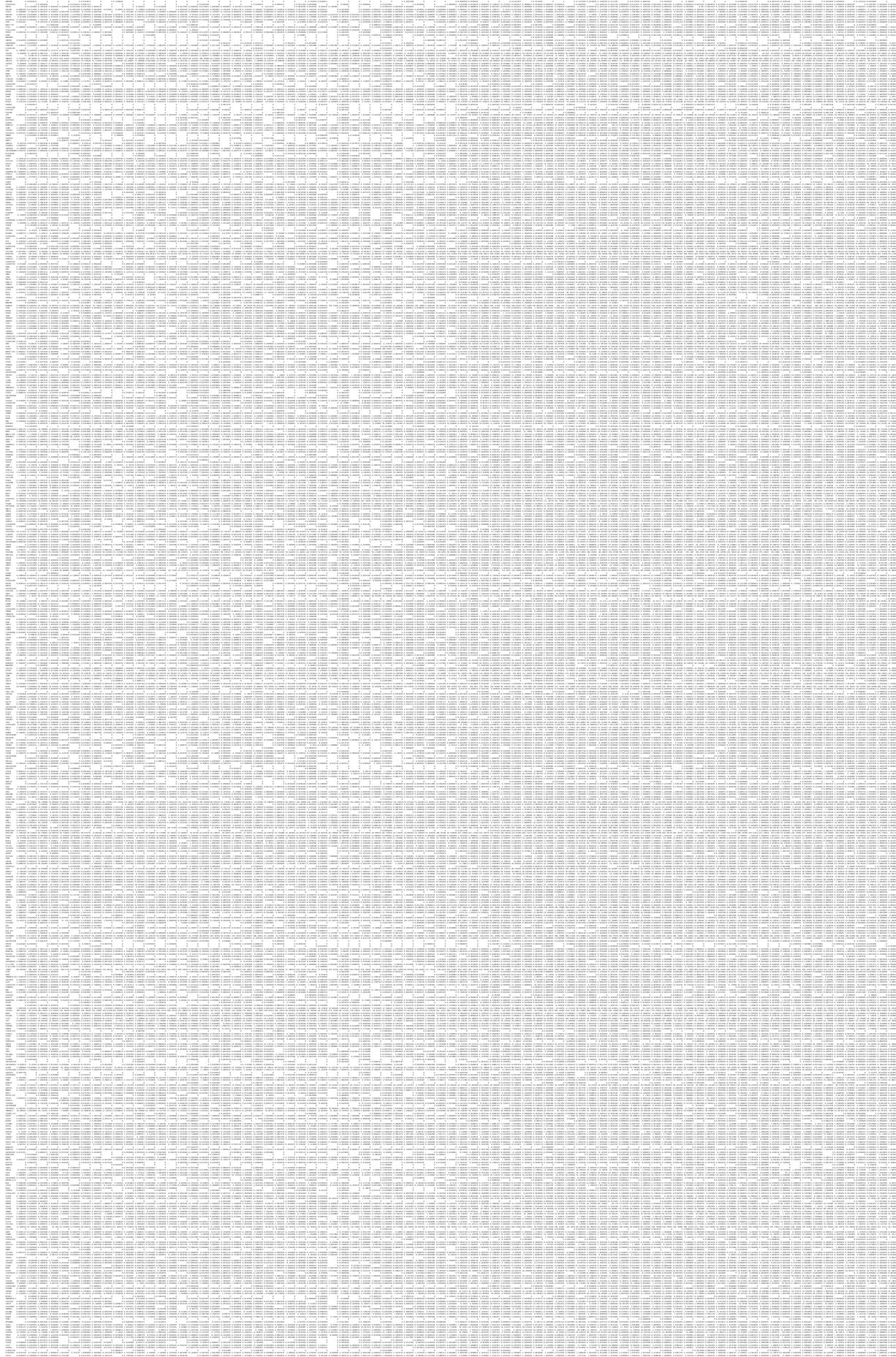
osteoporosis;Papillomavirus infection;Ovary tumor;Prostate tumor;Adrenal disease;Urinary dysfunction;Estrogen deficiency;Endometriosis;Hormone deficiency;Musculoskeletal disorder;Cardiovascular disease;Metabolic disorder;Neurological disorder;Gastrointestinal disorder;Hematological disorder;Infectious disease;Endocrine disorder;Genetic disorder;Immunological disorder;Musculoskeletal disorder;Cardiovascular disorder;Metabolic disorder;Neurological disorder;Gastrointestinal disorder;Hematological disorder;Infectious disease;Endocrine disorder;Genetic disorder;Immunological disorder

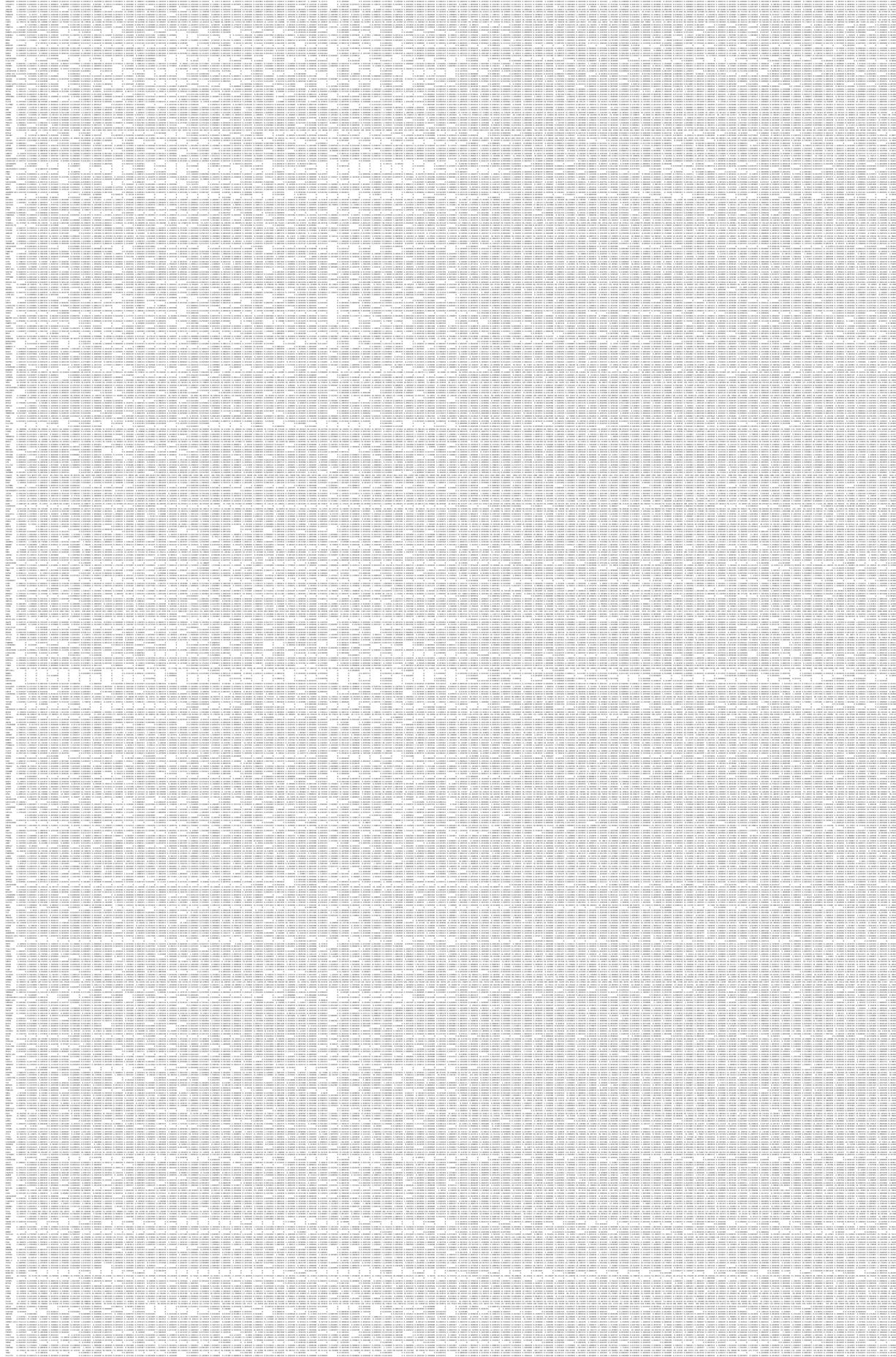
dystonia;Anemia;Endocrine disease;Non-insulin dependent diabetes;Breast tumor;Female contraception;Fungal infection;Cystic fibrosis;Prostate tumor;Estrogen deficiency;Endometriosis;Hormone deficiency;Musculoskeletal disorder;Cardiovascular disease;Metabolic disorder;Neurological disorder;Gastrointestinal disorder;Hematological disorder;Infectious disease;Endocrine disorder;Genetic disorder;Immunological disorder;Musculoskeletal disorder;Cardiovascular disorder;Metabolic disorder;Neurological disorder;Gastrointestinal disorder;Hematological disorder;Infectious disease;Endocrine disorder;Genetic disorder;Immunological disorder

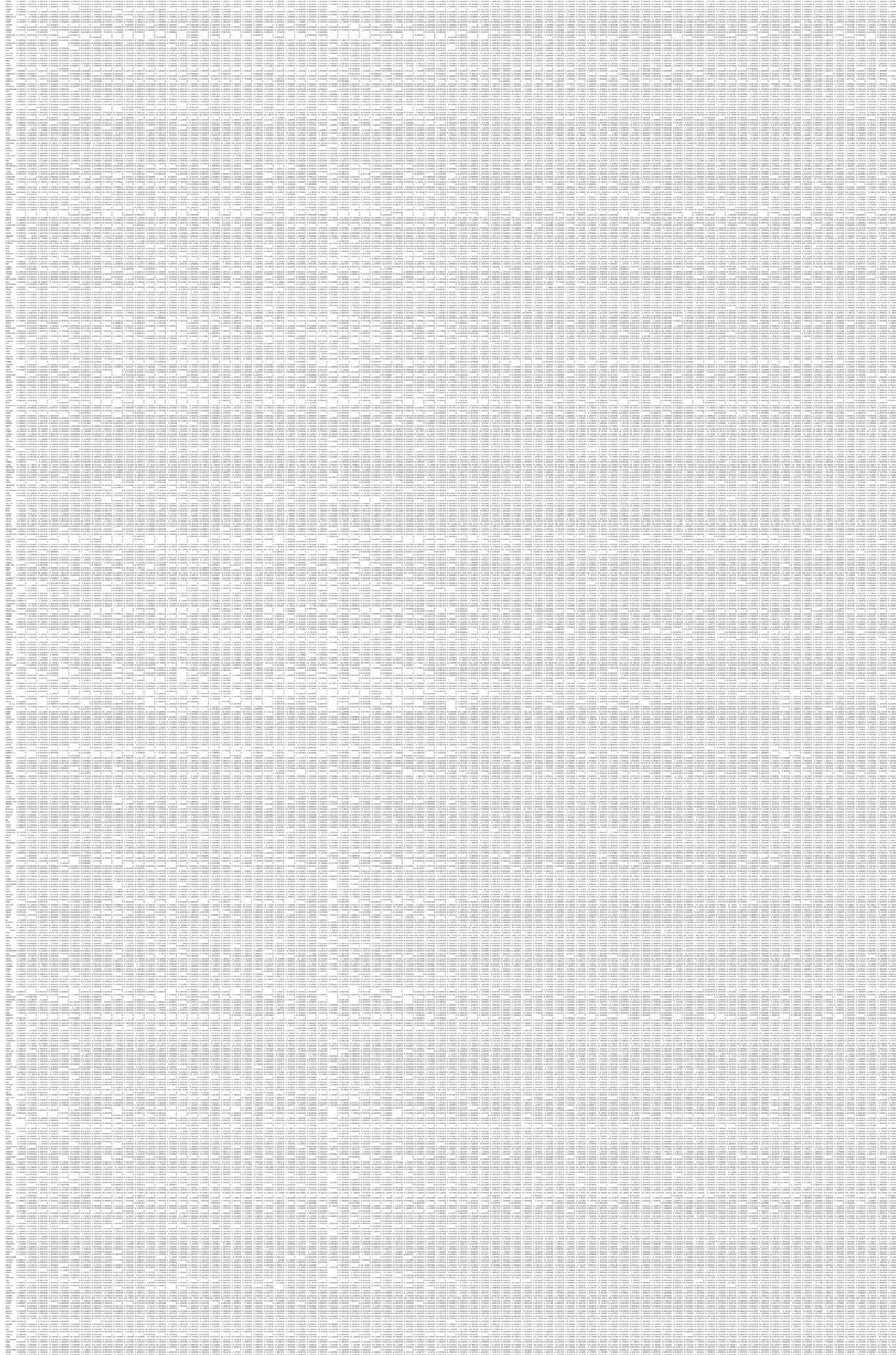
skeletal disease;Glioma;Vagina disease;Non-small-cell lung cancer;Parkinsons disease;Uterus tumor;Psychiatric disorder

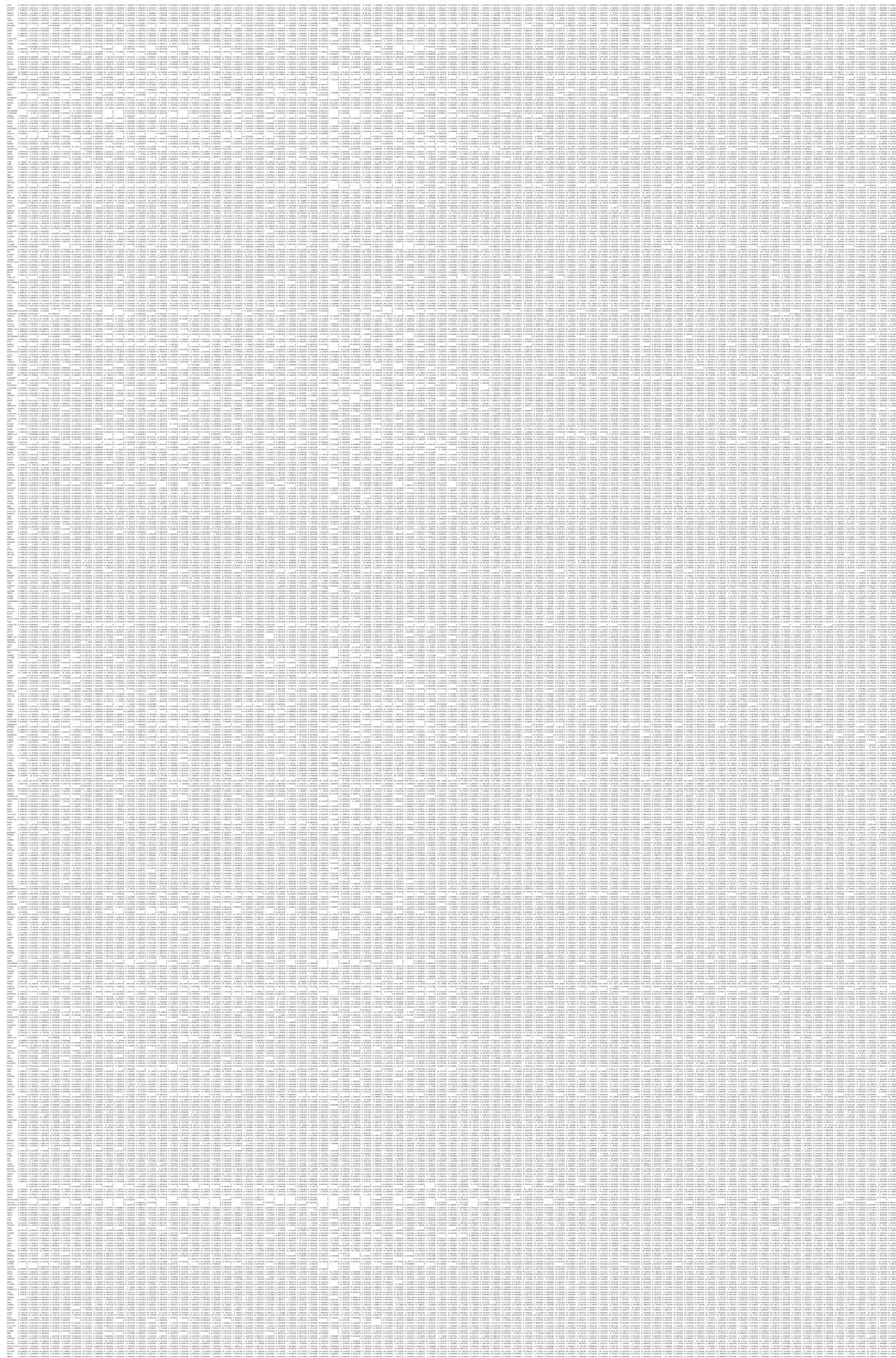
ency;Endometriosis;Hormone deficiency;Turners syndrome;Keratosis;Premenstrual syndrome;Hepatitis C virus infection;Benign tumor;Osteoporosis;Male contraception

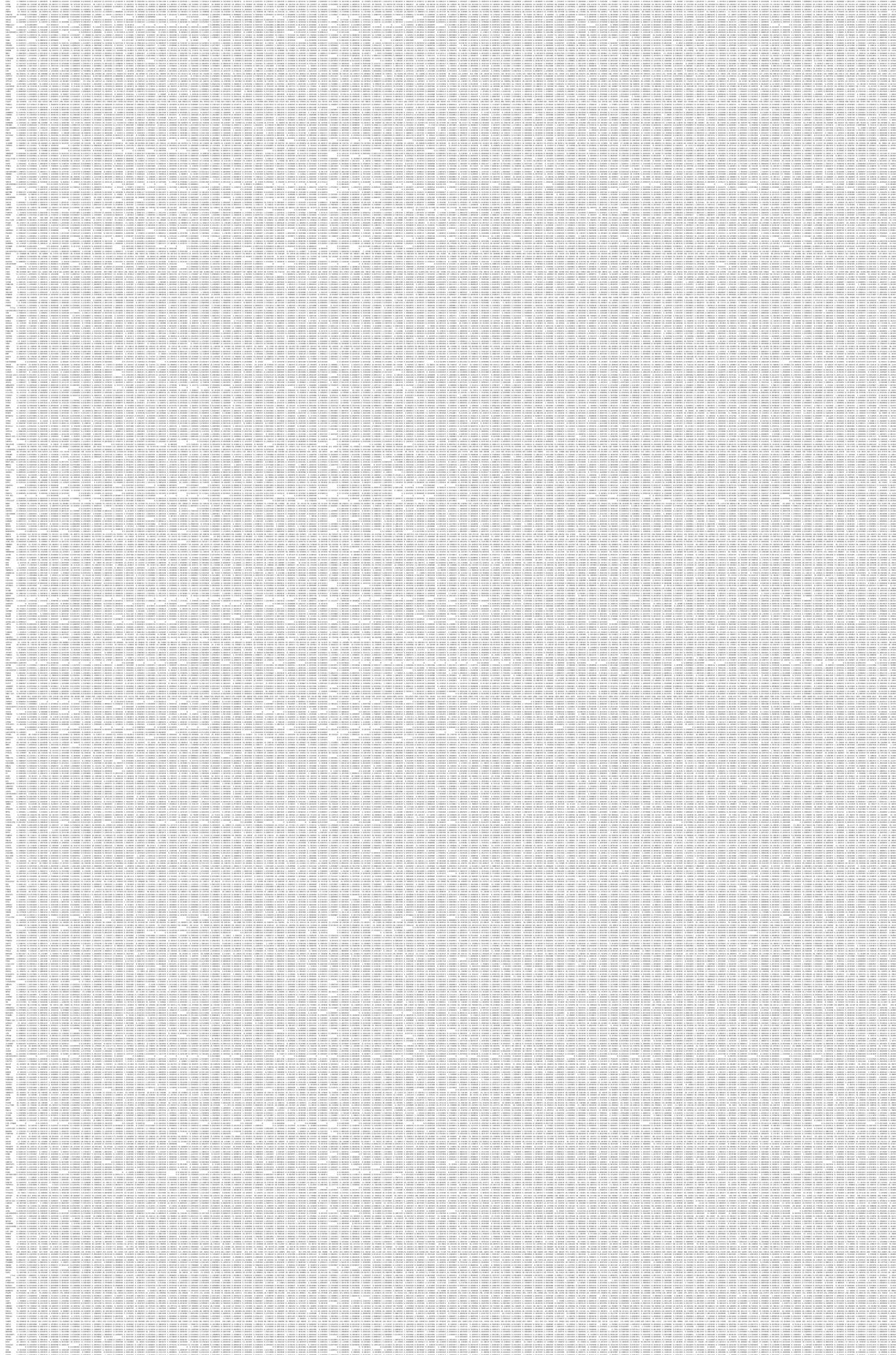


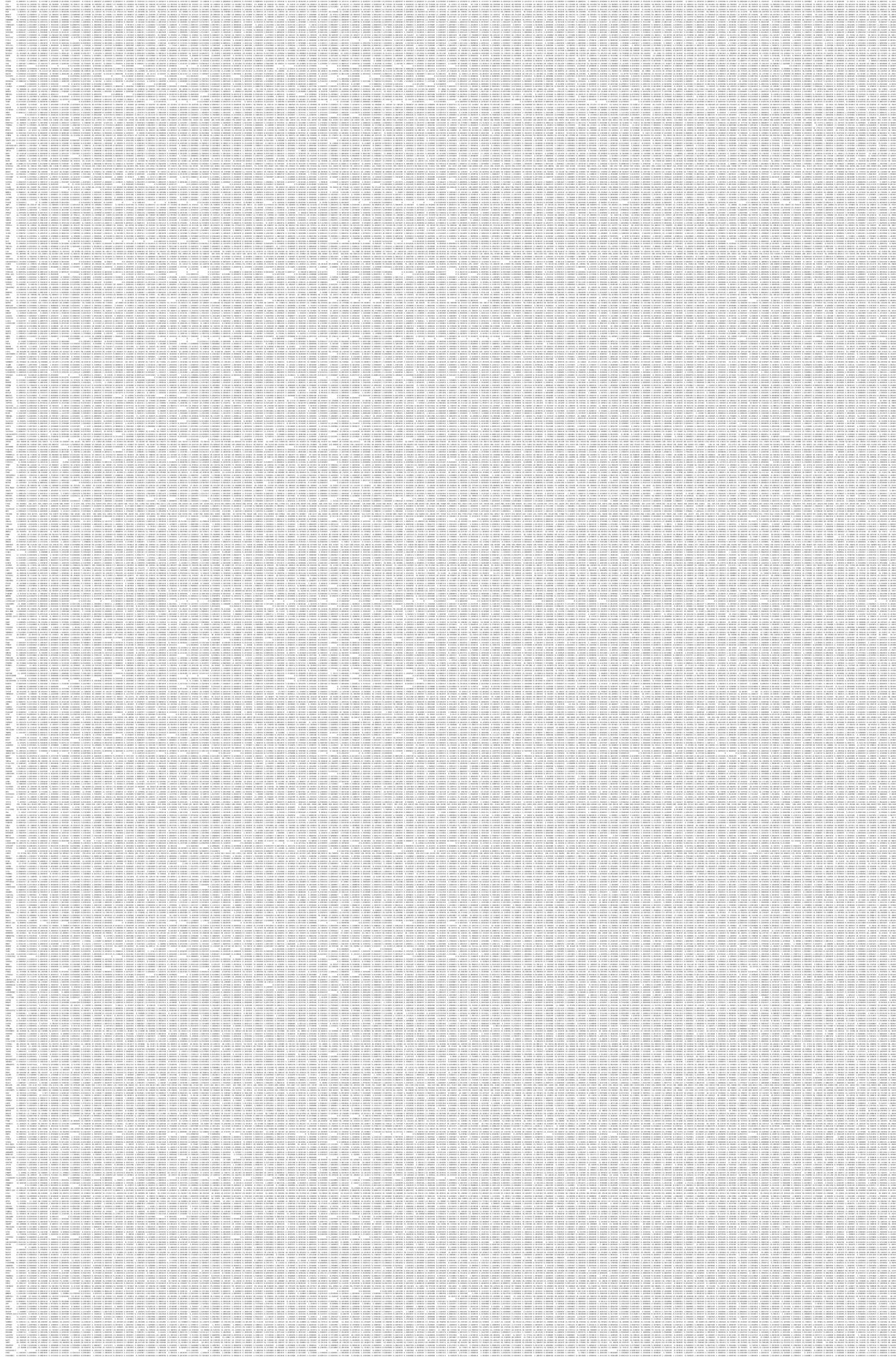


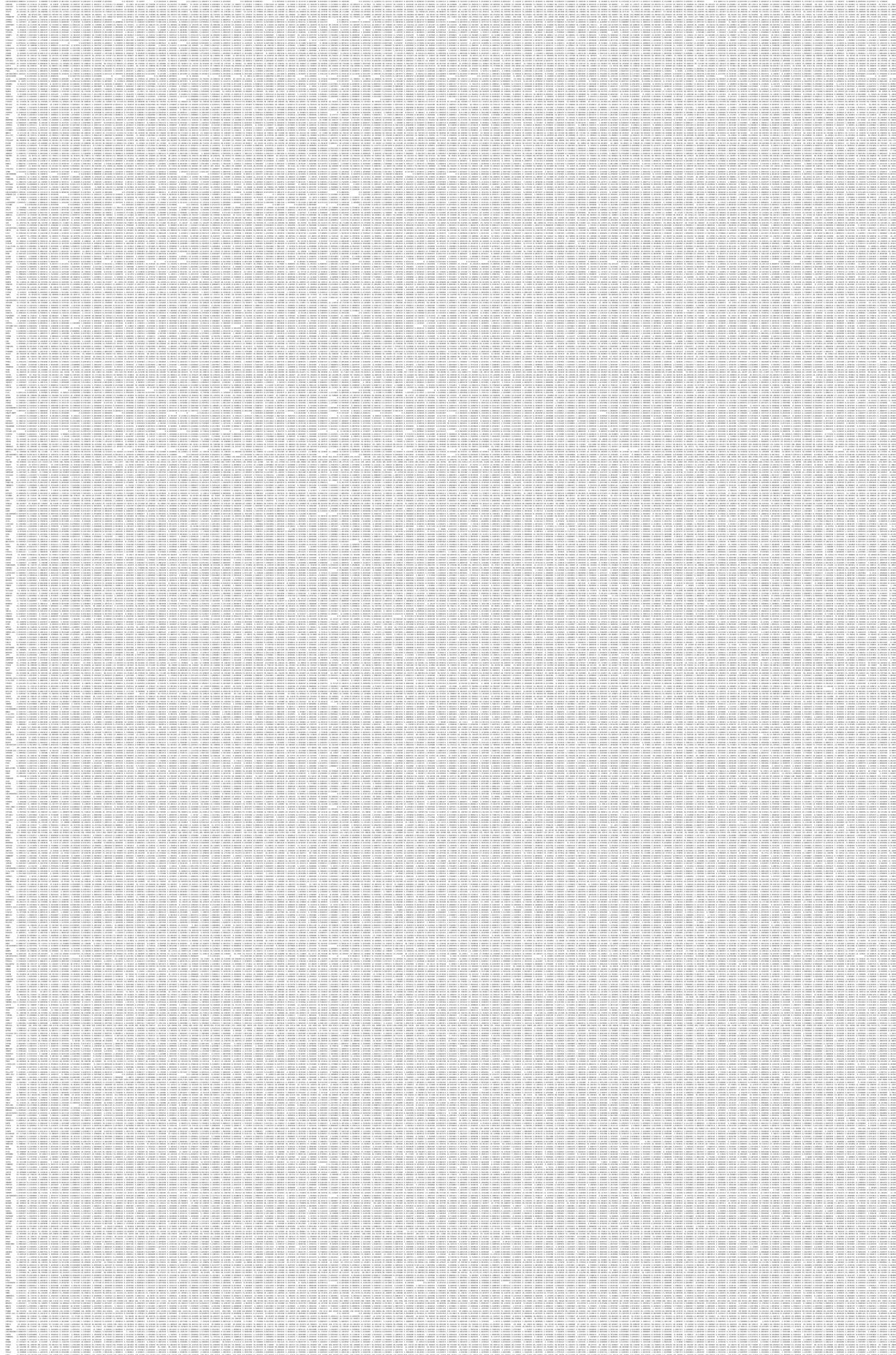


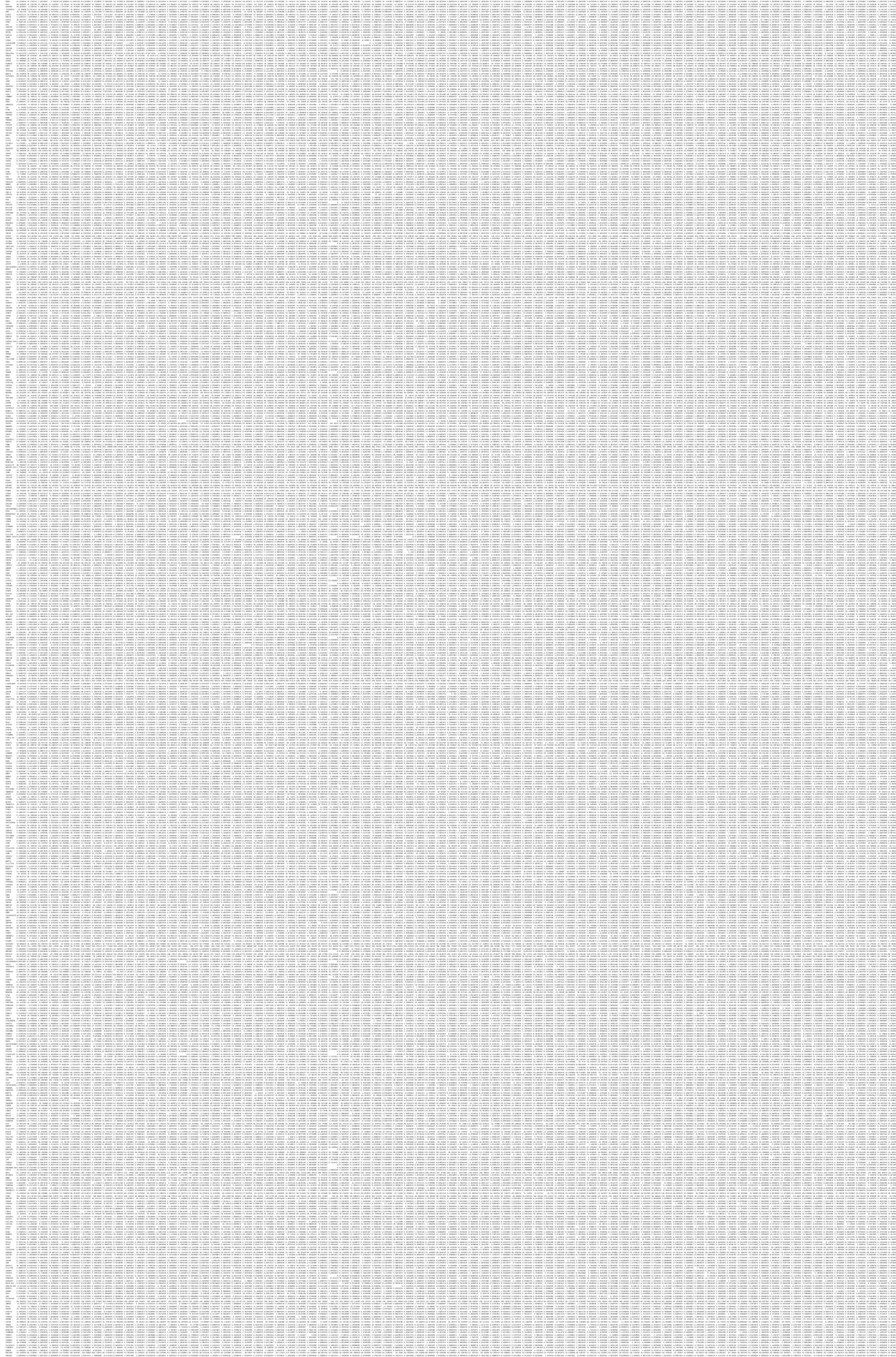


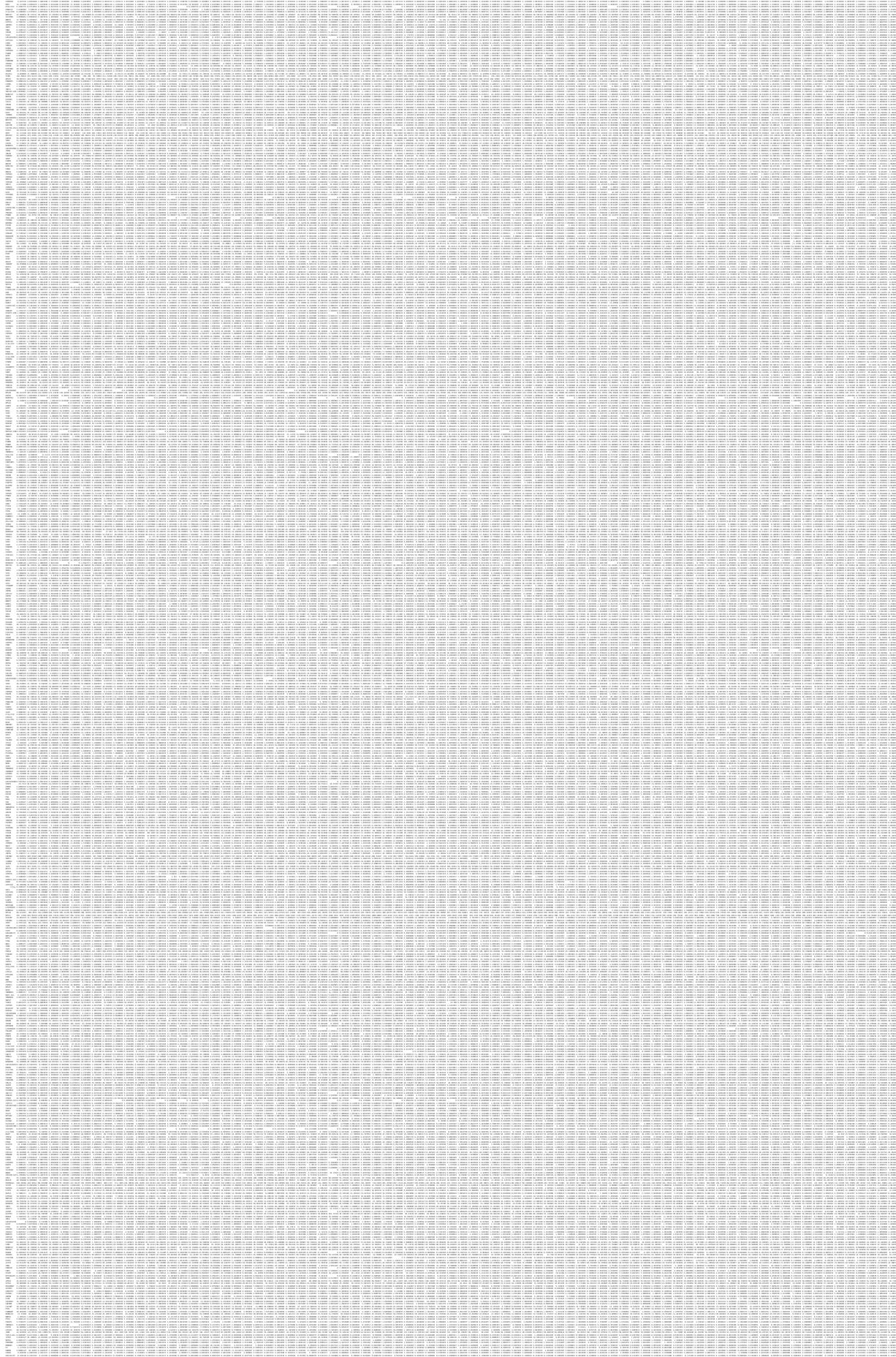


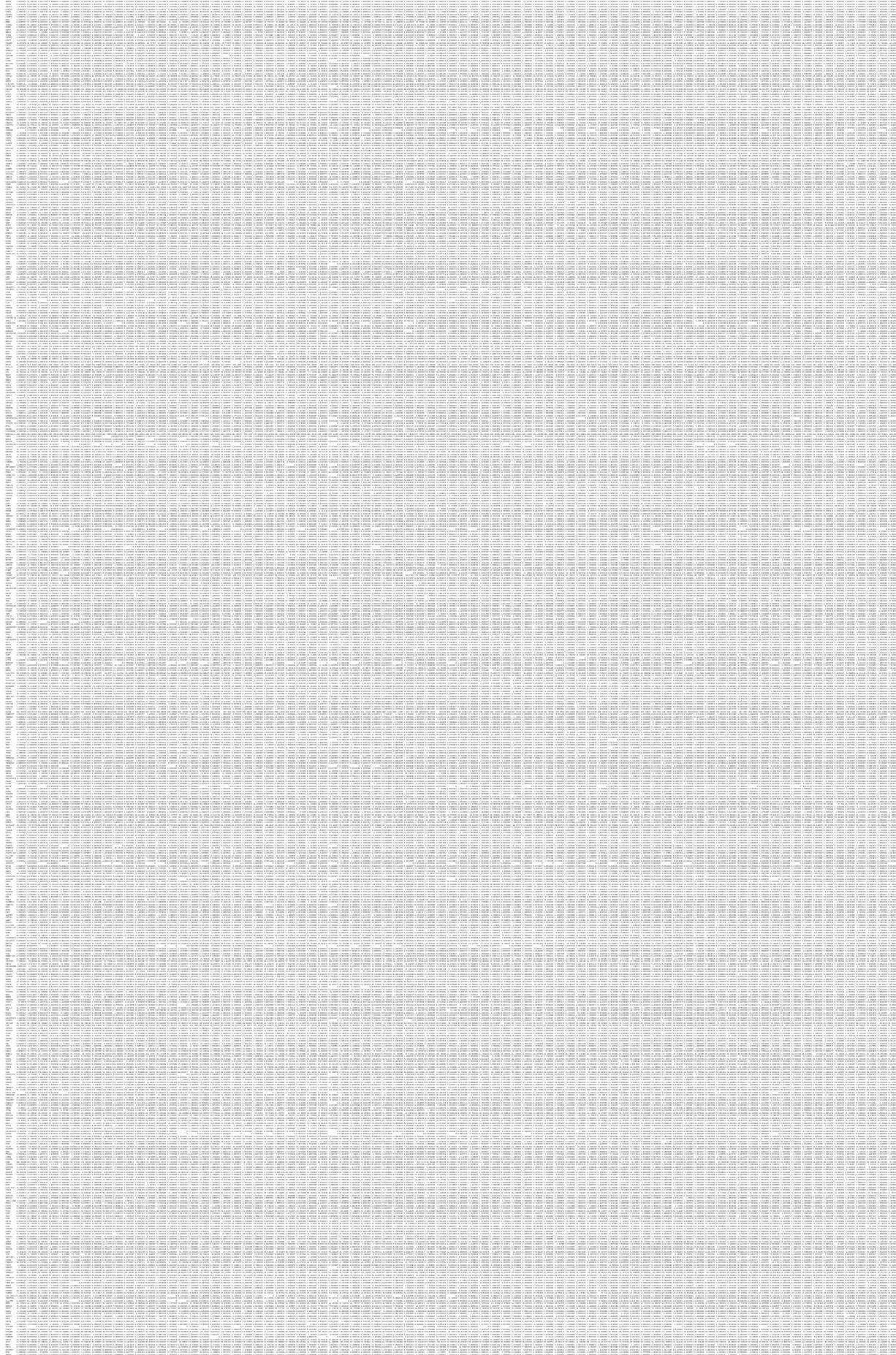


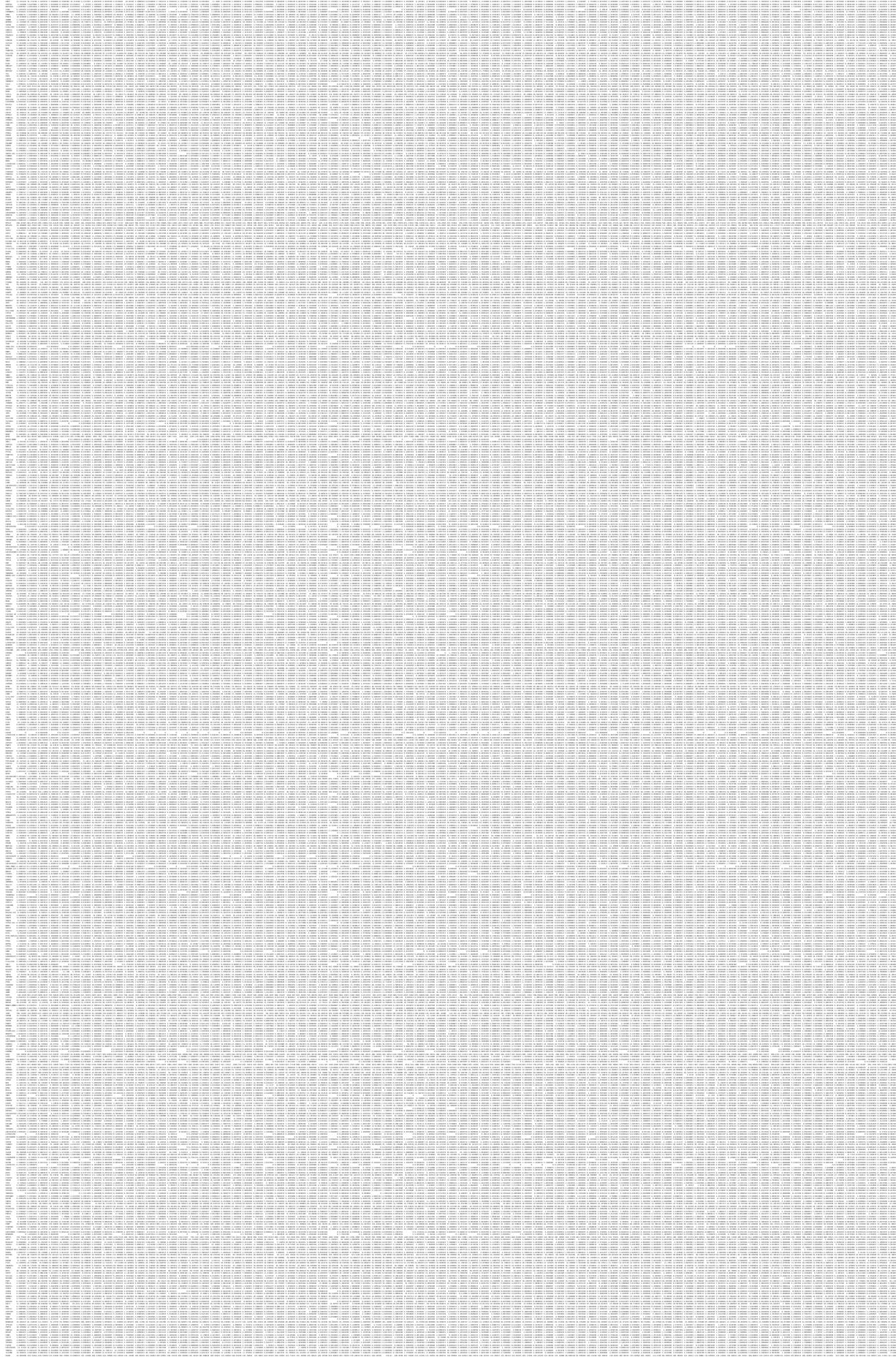


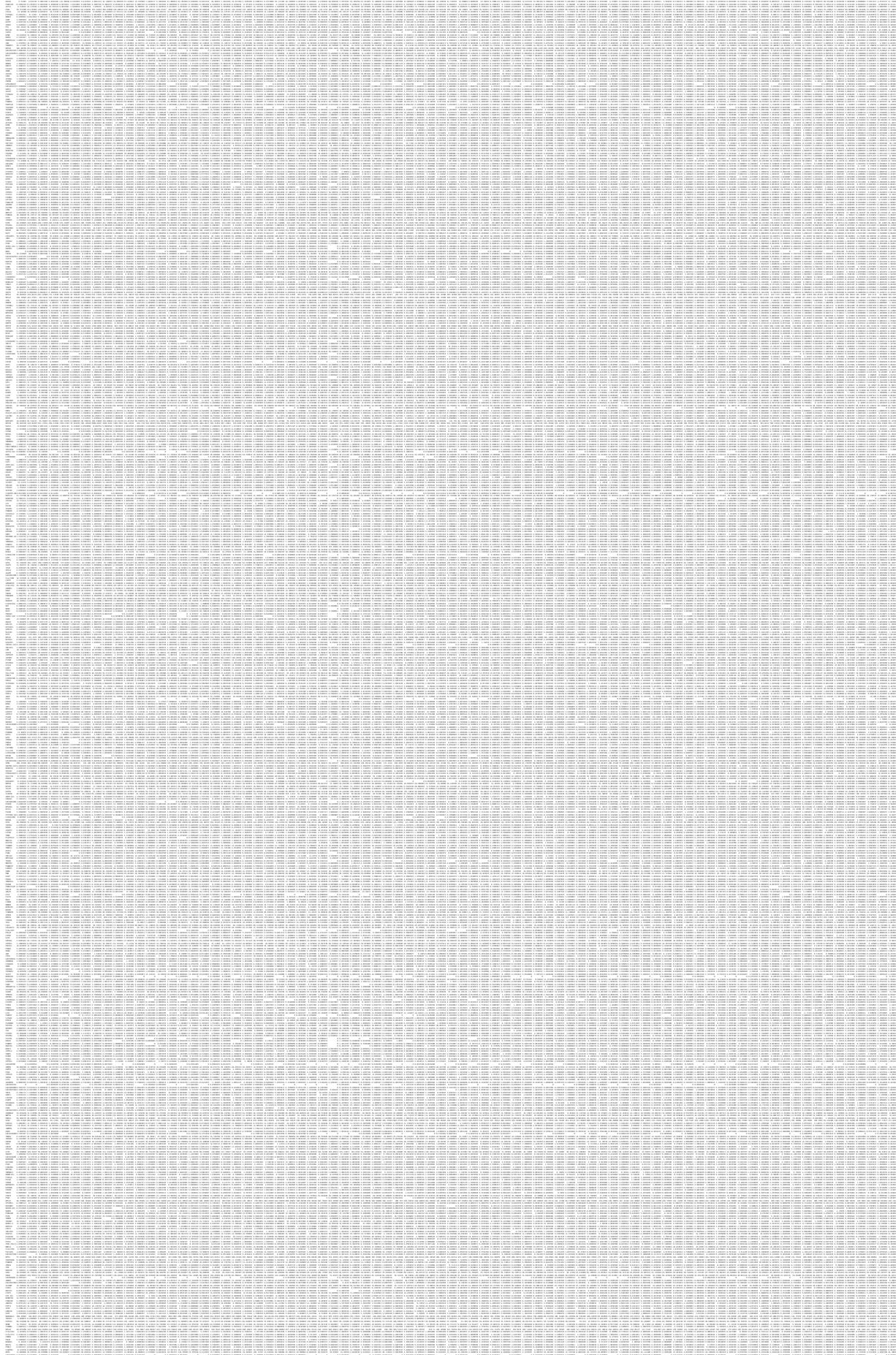


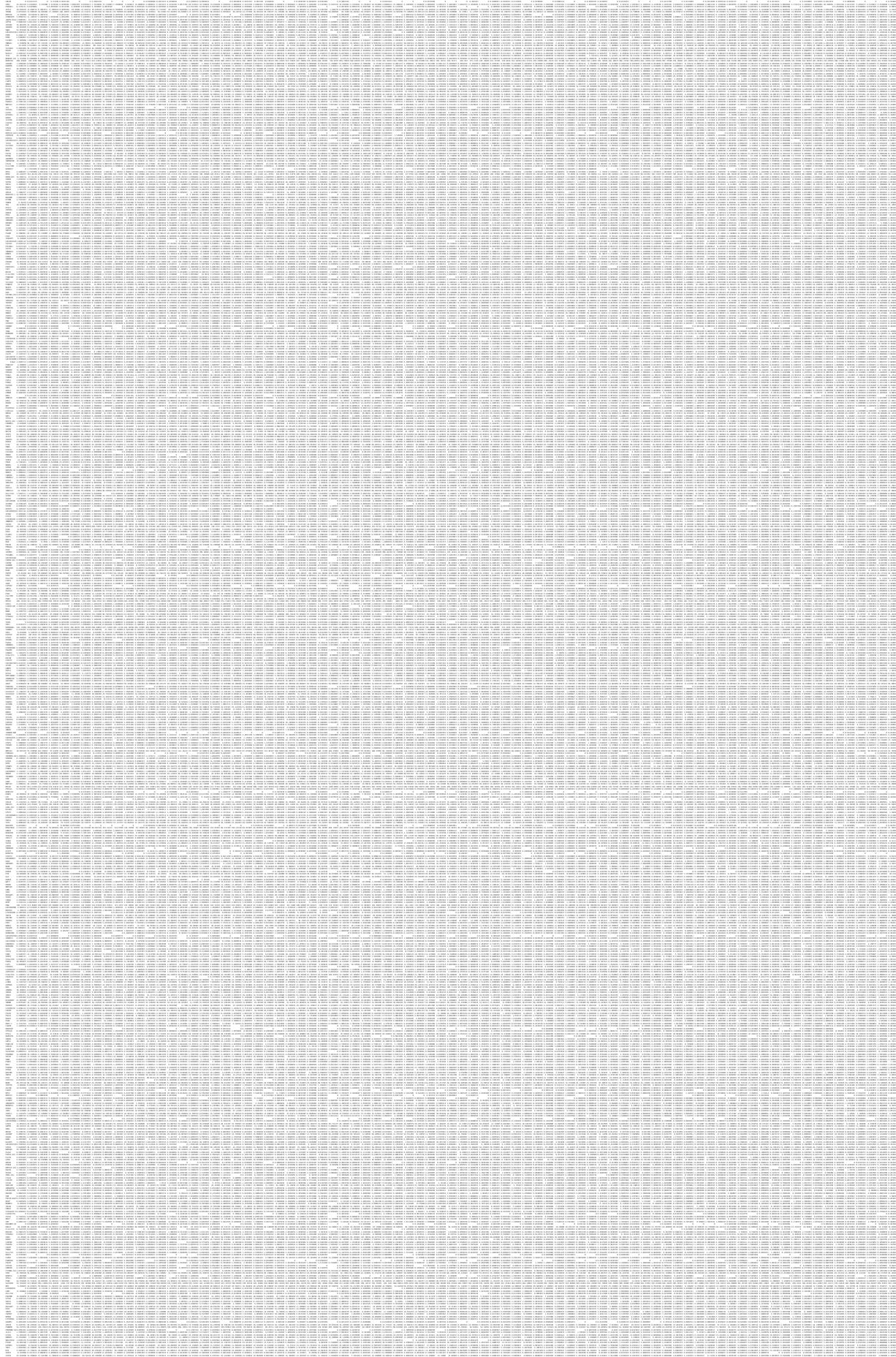


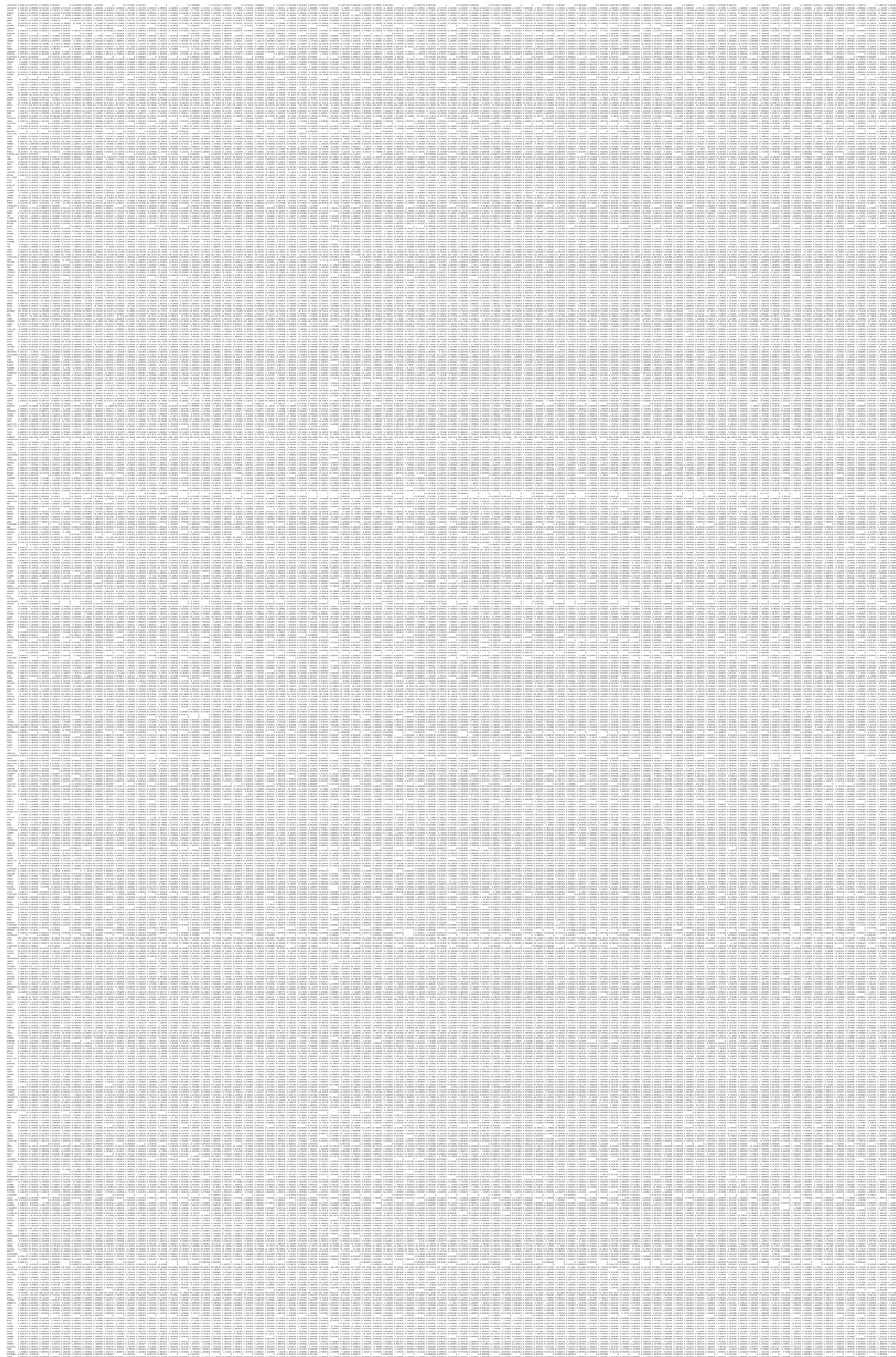


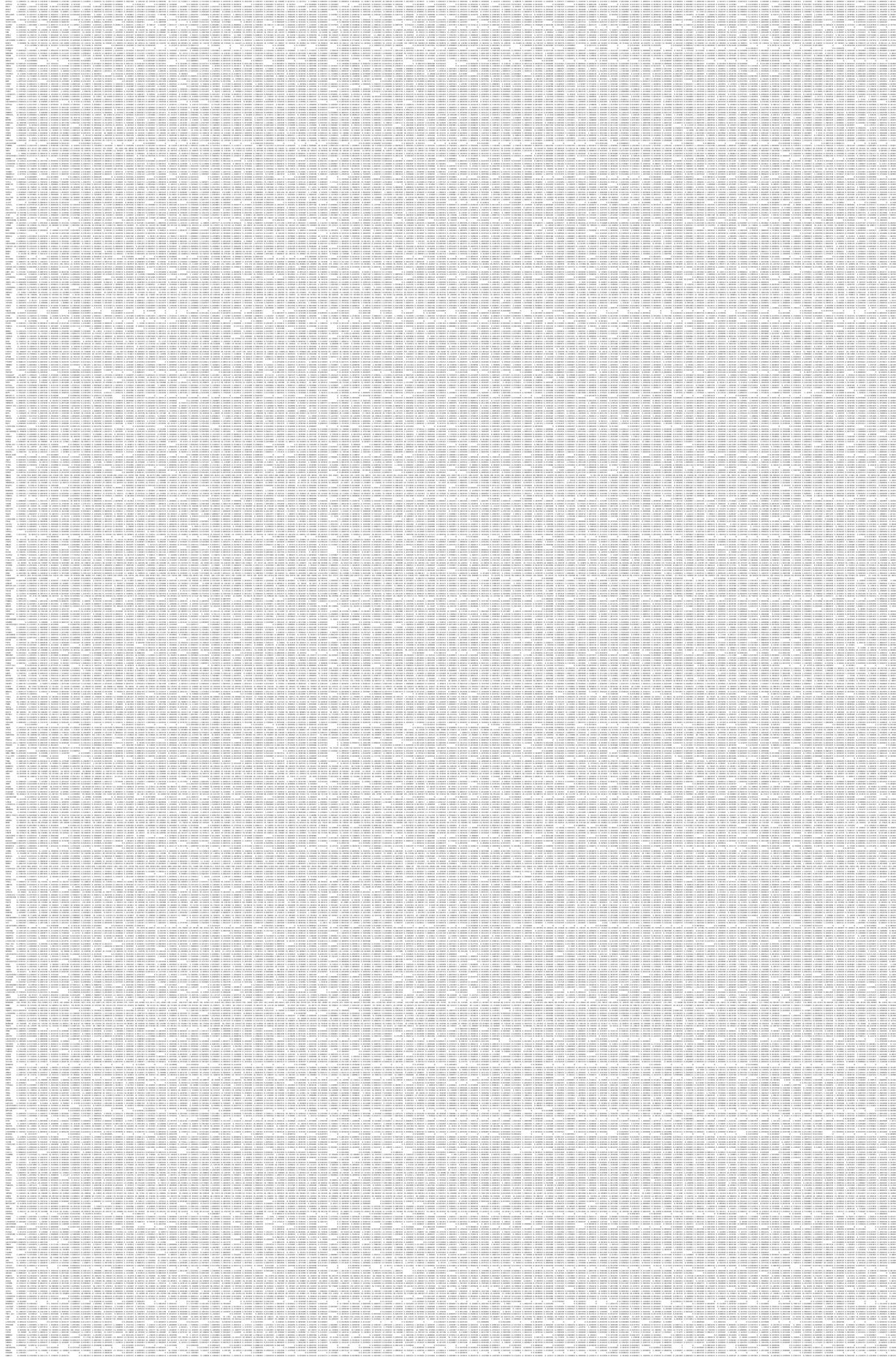


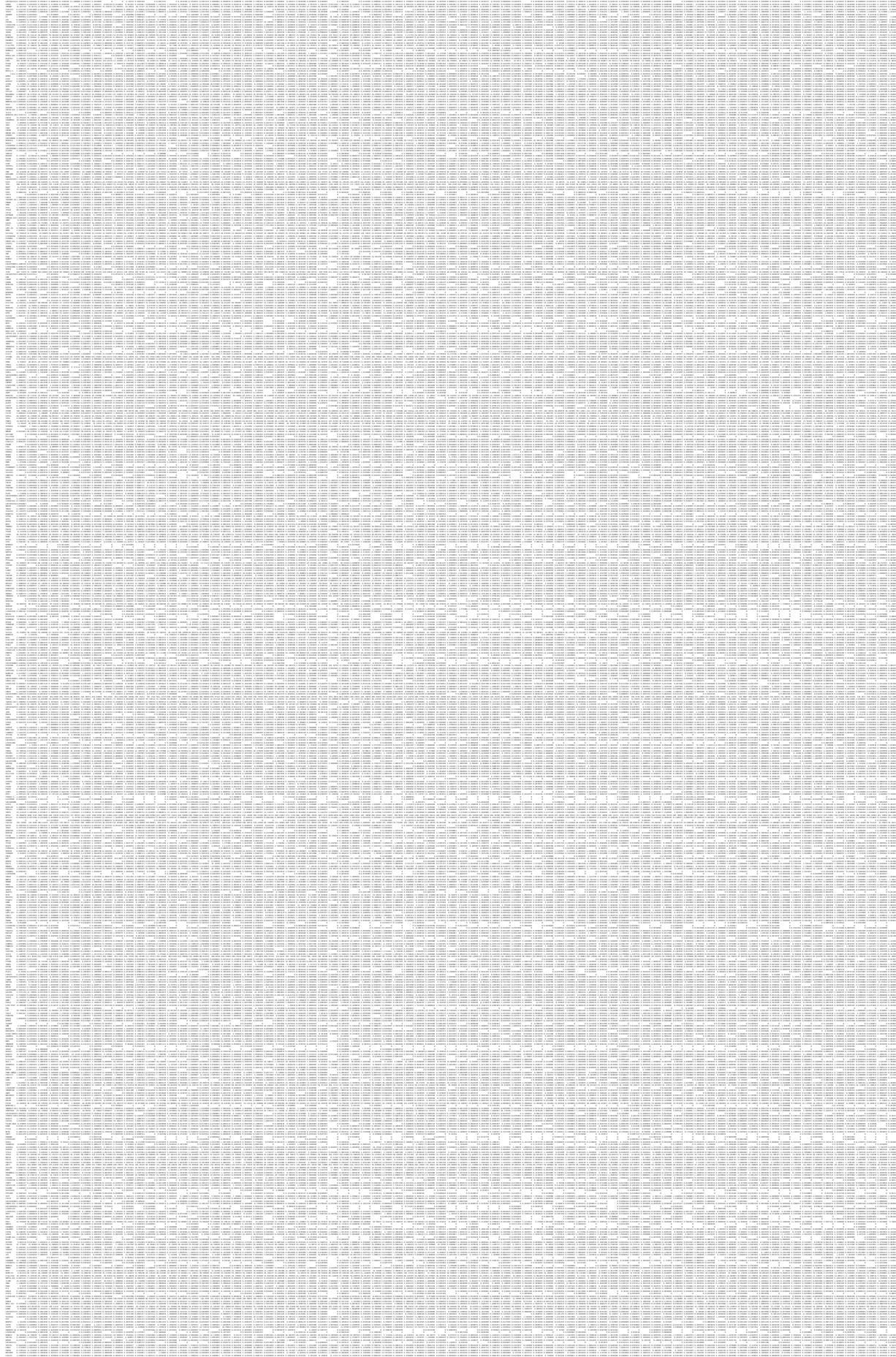


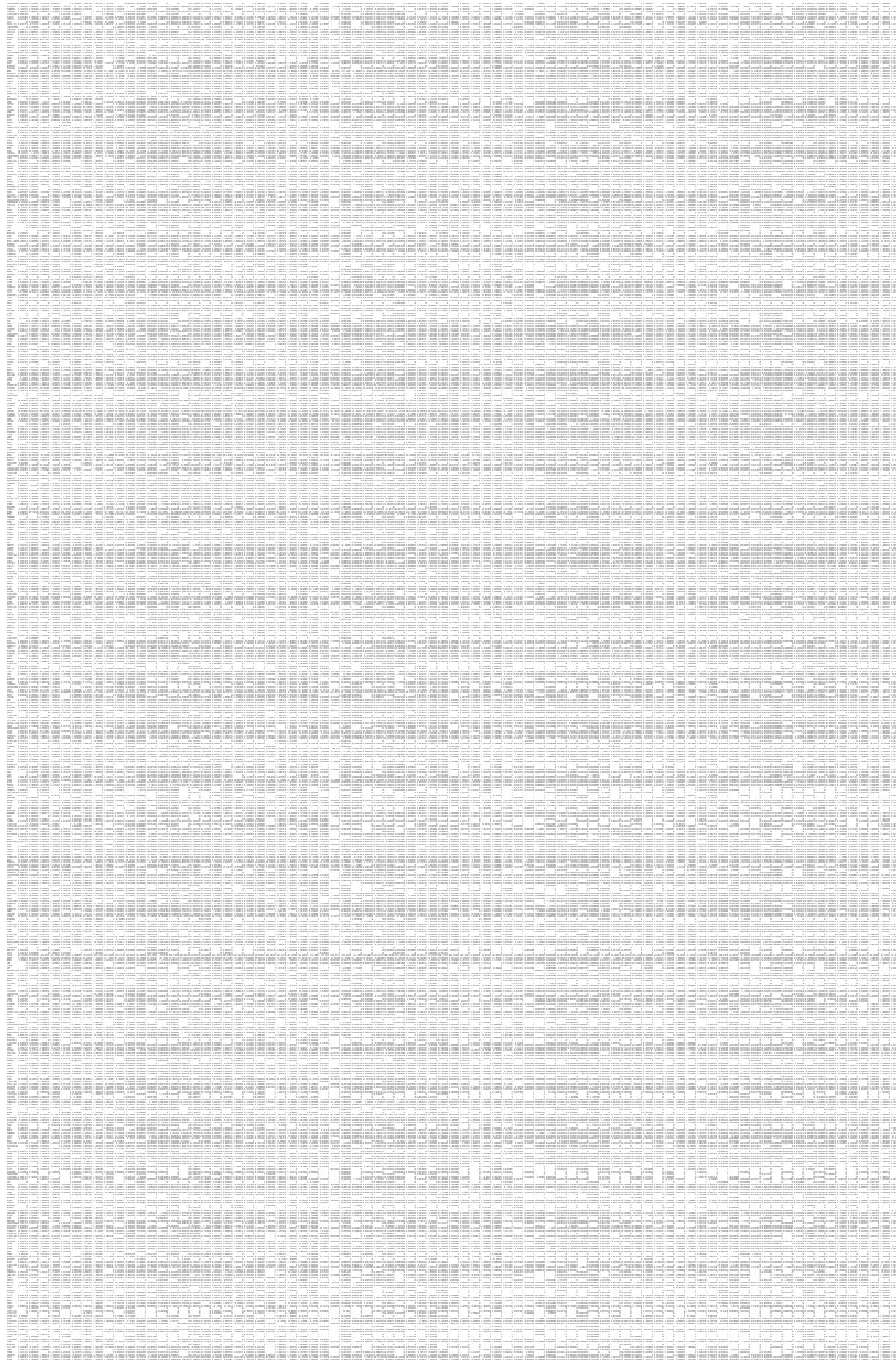












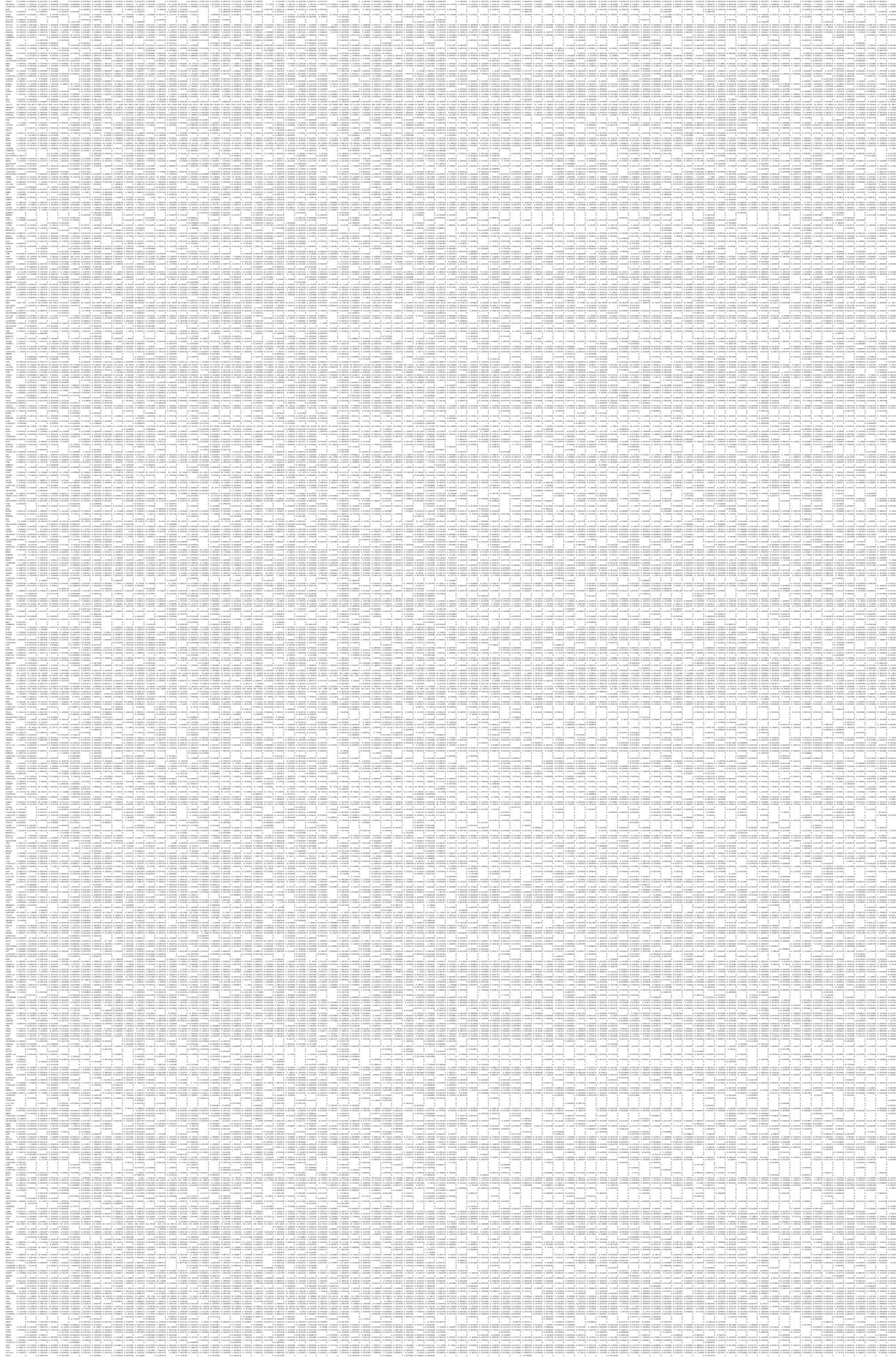




Table 4S

| List 1 | List 2 | Intersection |
|-------------------|----------|--------------|
| OTOP2 | MAPKAPK2 | CA1 |
| OTOP3 | TTR | CA2 |
| AQP8 | ALB | NR1H4 |
| KRT24 | STS | BCHE |
| CA7 | APOA2 | ADH1B |
| KRTAP13-2 | GC | TPH1 |
| MIR122 | ANG | ADH1C |
| SST | CASP3 | CLEC4M |
| TMIGD1 | PIM1 | PCK1 |
| BEST4 | BACE1 | CMA1 |
| DAO | AKR1B1 | ELANE |
| MYOC | #N/A | CPB1 |
| PYY | PDE4B | CTSG |
| PNLIPRP3 | GSR | NR3C2 |
| GUCA2B | MAPK1 | MAOB |
| MS4A12 | MAPK8 | CFD |
| CCDC169-SCSULT2A1 | | PGR |
| SLC30A10 | CES1 | THRB |
| PCSK2 | MAPK14 | LGALS2 |
| HSD3B2 | HSD17B1 | CD209 |
| LCE2A | AMPM2 | ERBB4 |
| KRT25 | TGFBR1 | IL2 |
| GCG | SHBG | KIT |
| DHRS7C | RORA | AKR1C2 |
| CD177 | LSS | BMP2 |
| LINC01289 | SRC | CYP2C8 |
| LINC00307 | FABP4 | FGFR2 |
| CLCA4 | DUSP6 | CASP7 |
| CHAT | REN | MAPK10 |
| SLC17A8 | CHIT1 | TPSB2 |
| ABCB11 | PTPN1 | PDE5A |
| LGI1 | HSD11B1 | FKBP1B |
| PABPC1L2B | ADH5 | PIK3CG |
| LINC00682 | KDR | NR3C1 |
| BMP3 | PTPN11 | KAT2B |
| PLP1 | PDPK1 | EPHX2 |
| SCNN1G | ESRRG | AR |
| ABCG2 | TNNC1 | HSD17B11 |
| LINC01445 | F10 | GSTM2 |
| SPIB | CSNK2A1 | AKR1C3 |
| RIT2 | TTPA | ACADM |
| PPY | PYGL | LGALS3 |
| HTR3E-AS1 | GSK3B | BST1 |
| CA4 | DPP4 | GP1BA |
| GUCA2A | PNMT | ITPKA |
| PRIMA1 | LCK | CTSS |
| OR8G5 | PDE3B | ITGAL |
| POU3F4 | PARP1 | CDA |
| PGM5-AS1 | AMY2A | CTSF |

| | | |
|-------------------|----------|----------|
| FRMPD4 | ISG20 | IMPA1 |
| DPP6 | LTA4H | FGFR1 |
| HS3ST6 | HMGCR | GSTM1 |
| NPTX1 | CYP2C9 | TEK |
| FUT9 | ALDH2 | VDR |
| SLC26A3 | SERPINA1 | PPARD |
| SLC6A19 | JAK3 | RXRA |
| LINC01976 | CTNNA1 | PDE4D |
| RBFOX3 | TRAPP C3 | PPARG |
| CWH43 | SOD2 | AKR1C1 |
| CLDN8 | GLRX | WAS |
| CEACAM7 | FABP7 | HNF4G |
| LINC01264 | REG1A | NMNAT1 |
| SERPINA9 | NR1H2 | ITK |
| RHAG | NR1H3 | EGFR |
| CMTM5 | PPARA | CBR1 |
| GBA3 | SULT1E1 | FECH |
| SLC4A4 | THRA | PDK2 |
| UGT1A8 | SYK | HCK |
| PHOX2B | GSTT2B | CASP1 |
| NRXN1 | GSTA1 | ESR1 |
| SCGN | XIAP | CDC42 |
| LINC00974 | FABP6 | ANXA5 |
| SCGB1D2 | FABP3 | MDM2 |
| CDKN2B-AS1PLA2G2A | | DHFR |
| NGB | RNASE4 | IGF1R |
| VSTM2A-OT1FHIT | | FDPS |
| TMEM72 | NR1I3 | PNP |
| SFRP5 | F7 | PPP5C |
| SCN7A | BLVRB | CTSB |
| VSTM2A | RHOA | ADAM17 |
| LINC01855 | AMY1B | AKT2 |
| STMN4 | FABP5 | PAPSS1 |
| ALPI | SELP | TYMS |
| SPHKAP | AMD1 | SDS |
| ZG16 | TPI1 | APRT |
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| GPM6A | NOS2 | GL01 |
| SLC26A2 | GM2A | CDK7 |
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| ATP1A2 | RARB | HSPA8 |
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| | | |
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| DCAF12L1 | MAN1B1 | EPHA2 |
| SLC51B | NT5M | CLPP |
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| GABRG2 | CSK | HDAC8 |
| CNTFR | AGXT | GART |
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| USP2 | HNMT | S100A9 |
| TRPM6 | PPCDC | GNPDA1 |
| SLC5A7 | HAGH | RARG |
| SLC51A | RXRB | NQO1 |
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| MORN5 | CCBL1 | DTYMK |
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| TMEFF2 | F11 | NQO2 |
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| SLC7A14 | PCTP | CCNA2 |
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| PI16 | CLK1 | AURKA |
| PABPC1L2A | PIK3R1 | EPHB4 |
| FEV | GCDH | RBP4 |
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| MYH11 | TRDMT1 | IMPDH1 |
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| TCEAL2 | | MMP12 |
| MMP27 | | AHCY |
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1-Sep

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4-Mar

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LINC01657
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STC2
SRPX2
ANOS1
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FAP
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HOXC-AS2
TRIM54
MEPE
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HCAR1
DLX6
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PLEKHN1
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LYZL4
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SPP1
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CHST4
SPZ1
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HOXC11

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LINC01977
ADAM12
ADAM18
VWA2
CALCA
TG
SLC22A12
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DMRTA2
GJB5
ULBP2
TCF24
EPHA8
KRT17
UNC93A
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IRX5
IL11
SLC13A3
VGLL1
SIM2
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FEZF2
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CCAT2
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CLDN2
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MAP3K20-AS1
OFCC1
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KRT83
TACSTD2
MIR566
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SPRR3
AKAP4
SALL4
EIF3IP1
GRIN2D
UCN2
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TCAM1P
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CXCL17
ELFN1-AS1

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ETV4
OTOP1
GAD1
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RP1
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NEUROG2
SNHG25
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LINC00534
KLK10
PIWIL1
INHBA
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ESM1
KRT38
ZIC2
AKR1C4
SERPINA4
FOXI3
BOD1L2
AMELX
LINC02119
WNT7B
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CLDN10-AS1
MIR613
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FAR2P1
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LEMD1
FIRRE

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KLC3
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SPRR1B
C3P1
NXPH4
CPNE7
MIR135B
PCAT2
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GBX2
ONECUT3
FAM9A
SPRR2A
CRAT37
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LOC284930
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MIR4474
STRA6
LINC01411
LY6G6F
OBP2B
MAGEA11
IFITM5
KRT35
SLC01B3
C14orf105
KRT32
SPRR2D
FOXQ1
LINC01169
AFAP1-AS1
C5orf46
COL11A1
IBSP
BMS1P22
MMP7
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LOC101929448
KRT80
CST2
SPRR1A
LINC01655
CEL
LOC102724908
LINC01913

KRT23
SNORD30
LINC02187
SFTA2
PTF1A
EPYC
DUSP27
ELF5
LINC01602
SPERT
LINC01050
SOX14
LINC01234
COL10A1
ZIC5
LINC01593
C6orf15
CST4
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KLK7
KLK8
FEZF1
KLK6
GOLGA6L2
FEZF1-AS1
LOC100190940
RNA5S9
NOTUM

Table 5S

| node1 | node2 | node1_strin | node2_strin | node1_extenode2 | exteneighbor | homologe_fusion | coexpressi | experiment | database | _automated_ | combined_score | | |
|----------|----------|-------------|-------------|------------------------|--------------|-----------------|------------|------------|----------|-------------|----------------|-------|-------|
| CDC42 | WAS | 4446990 | 4444767 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.063 | 0.974 | 0.9 | 0.97 | 0.999 |
| HSPA8 | HSP90AA1 | 4449616 | 4440643 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.771 | 0.73 | 0.9 | 0.929 | 0.999 |
| DHFR | TYMS | 4447922 | 4439011 | 9606. ENSP09606. ENSP0 | 0.219 | 0.317 | 0.437 | 0 | 0.479 | 0.122 | 0.9 | 0.96 | 0.999 |
| CCNA2 | CDK2 | 4436109 | 4435732 | 9606. ENSP09606. ENSP0 | 0 | 0.006 | 0 | 0 | 0.403 | 0.998 | 0.9 | 0.973 | 0.999 |
| CCNA2 | CDK6 | 4436109 | 4435642 | 9606. ENSP09606. ENSP0 | 0 | 0.003 | 0 | 0 | 0.163 | 0.994 | 0.72 | 0.769 | 0.999 |
| CDC42 | TNK2 | 4446990 | 4445619 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.974 | 0.9 | 0.641 | 0.998 |
| HSPA8 | HSP90AB1 | 4449616 | 4433770 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.641 | 0.745 | 0.9 | 0.815 | 0.998 |
| RXRA | PPARG | 4449061 | 4436745 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0.694 | 0.061 | 0.961 | 0.9 | 0.821 | 0.997 |
| TYMS | TK1 | 4439011 | 4437779 | 9606. ENSP09606. ENSP0 | 0.169 | 0 | 0 | 0 | 0.787 | 0 | 0.9 | 0.856 | 0.997 |
| HSP90AB1 | HSP90AA1 | 4443770 | 4440643 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0.449 | 0.984 | 0.78 | 0.883 | 0.9 | 0.717 | 0.997 |
| GART | MTHFD1 | 4445605 | 4432975 | 9606. ENSP09606. ENSP0 | 0.37 | 0 | 0 | 0 | 0.695 | 0 | 0.9 | 0.82 | 0.996 |
| AKR1C1 | AKR1C3 | 4445467 | 4445414 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0.449 | 0.983 | 0.214 | 0.943 | 0.9 | 0.836 | 0.995 |
| RXRA | THR8 | 4449061 | 4446582 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0.691 | 0 | 0.974 | 0.8 | 0.782 | 0.995 |
| KIF11 | AURKA | 4434801 | 4432988 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.94 | 0.482 | 0 | 0.831 | 0.994 |
| GP1BA | F2 | 4440098 | 4438421 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.909 | 0.9 | 0.481 | 0.994 |
| TYMS | DTYMK | 4439011 | 4438077 | 9606. ENSP09606. ENSP0 | 0.182 | 0 | 0 | 0 | 0.467 | 0.05 | 0.9 | 0.839 | 0.992 |
| NR1H4 | RXRA | 4450225 | 4449061 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0.668 | 0.062 | 0.904 | 0.9 | 0.745 | 0.992 |
| APRT | ADK | 4445041 | 4436683 | 9606. ENSP09606. ENSP0 | 0.043 | 0 | 0 | 0 | 0.088 | 0.159 | 0.9 | 0.909 | 0.992 |
| HSP90AA1 | NOS3 | 4440643 | 4437409 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.06 | 0.379 | 0.9 | 0.873 | 0.991 |
| DTYMK | TK1 | 4438077 | 4437779 | 9606. ENSP09606. ENSP0 | 0.05 | 0 | 0 | 0 | 0.569 | 0 | 0.9 | 0.817 | 0.991 |
| AKR1C1 | AKR1C2 | 4445467 | 4445450 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0.449 | 0.986 | 0.405 | 0.86 | 0.9 | 0.892 | 0.991 |
| CCNA2 | KIF11 | 4436109 | 4434801 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.977 | 0.15 | 0 | 0.478 | 0.989 |
| PNP | ADK | 4442485 | 4436683 | 9606. ENSP09606. ENSP0 | 0.167 | 0 | 0 | 0 | 0.061 | 0 | 0.9 | 0.88 | 0.989 |
| ESR1 | IGF1R | 4448358 | 4435797 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.407 | 0.9 | 0.835 | 0.989 |
| HSP90AA1 | EGFR | 4440643 | 4436186 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.487 | 0.9 | 0.817 | 0.989 |
| HSP90AA1 | PPP5C | 4440643 | 4432591 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.106 | 0.581 | 0.9 | 0.718 | 0.988 |
| AR | HSP90AA1 | 4444436 | 4440643 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.474 | 0.9 | 0.803 | 0.988 |
| DHFR | GART | 4447922 | 4456063 | 9606. ENSP09606. ENSP0 | 0.346 | 0 | 0 | 0 | 0.116 | 0 | 0.9 | 0.827 | 0.988 |
| RXRA | RARG | 4449061 | 4447554 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0.79 | 0.084 | 0.853 | 0.9 | 0.759 | 0.987 |
| CCNA2 | AURKA | 4436109 | 4432988 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.964 | 0.177 | 0 | 0.582 | 0.986 |
| HSP90AA1 | PGR | 4440643 | 4439740 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.523 | 0.9 | 0.72 | 0.985 |
| AKR1C2 | AKR1C3 | 4445450 | 4445414 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0.449 | 0.983 | 0.214 | 0.912 | 0.8 | 0.836 | 0.985 |
| ESR1 | NOS3 | 4448358 | 4437409 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.472 | 0.9 | 0.749 | 0.985 |
| APRT | PNP | 4445041 | 4442485 | 9606. ENSP09606. ENSP0 | 0.219 | 0 | 0 | 0 | 0.084 | 0 | 0.9 | 0.805 | 0.984 |
| TYMS | MTHFD1 | 4439011 | 4432975 | 9606. ENSP09606. ENSP0 | 0.208 | 0 | 0 | 0 | 0.159 | 0 | 0.9 | 0.79 | 0.984 |
| HSP90AA1 | NR3C1 | 4440643 | 4433539 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.474 | 0.9 | 0.71 | 0.983 |
| AKT2 | HSP90AA1 | 4445979 | 4440643 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.061 | 0.713 | 0.9 | 0.428 | 0.982 |
| ELANE | CTSG | 4450799 | 4432948 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0.437 | 0.821 | 0.784 | 0 | 0.9 | 0.92 | 0.982 |
| CCNA2 | CDK7 | 4436109 | 4434485 | 9606. ENSP09606. ENSP0 | 0 | 0.003 | 0 | 0 | 0.081 | 0.451 | 0.9 | 0.669 | 0.981 |
| CDA | TK1 | 4444499 | 4437779 | 9606. ENSP09606. ENSP0 | 0.292 | 0 | 0 | 0 | 0.061 | 0 | 0.9 | 0.742 | 0.98 |
| DHFR | MTHFD1 | 4447922 | 4432975 | 9606. ENSP09606. ENSP0 | 0.146 | 0 | 0 | 0 | 0.17 | 0 | 0.9 | 0.739 | 0.979 |
| ESR1 | HSP90AA1 | 4448358 | 4440643 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.474 | 0.9 | 0.633 | 0.979 |
| HDAC8 | KAT2B | 4444214 | 4435311 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.094 | 0.266 | 0.9 | 0.708 | 0.978 |
| AR | HSP90AB1 | 4444436 | 4437770 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.474 | 0.9 | 0.609 | 0.977 |
| HSP90AB1 | PPP5C | 4443770 | 4432591 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.103 | 0.581 | 0.9 | 0.456 | 0.976 |
| CDA | PNP | 4444499 | 4442485 | 9606. ENSP09606. ENSP0 | 0.219 | 0 | 0.215 | 0 | 0.077 | 0 | 0.9 | 0.642 | 0.976 |
| HCK | EGFR | 4450141 | 4436186 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0.679 | 0 | 0.701 | 0.9 | 0.722 | 0.975 |
| CDK2 | CDK6 | 4435732 | 4435642 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0.439 | 0.927 | 0.108 | 0.711 | 0.9 | 0.908 | 0.974 |
| ADH1C | ADH1B | 4449347 | 4438237 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0.449 | 0.986 | 0.179 | 0.692 | 0.9 | 0.921 | 0.973 |
| CDC42 | EGFR | 4446990 | 4436186 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.126 | 0.9 | 0.714 | 0.972 |
| ERBB4 | ADAM17 | 4441238 | 4438549 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.379 | 0.9 | 0.57 | 0.971 |
| DHFR | CBR1 | 4447922 | 4436857 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.062 | 0 | 0.9 | 0.725 | 0.971 |
| CDK2 | MDM2 | 4435732 | 4434604 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.05 | 0 | 0.9 | 0.697 | 0.968 |
| HSP90AB1 | NR3C1 | 4443770 | 4433539 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.403 | 0.9 | 0.509 | 0.968 |
| CHEK1 | MDM2 | 4447561 | 4434604 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0 | 0.9 | 0.698 | 0.968 |
| AR | CDK6 | 4444436 | 4435642 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.524 | 0.9 | 0.399 | 0.968 |
| ELANE | MMP8 | 4450799 | 4433667 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.175 | 0 | 0.9 | 0.633 | 0.967 |
| AR | KAT2B | 4444436 | 4435311 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.478 | 0.9 | 0.43 | 0.967 |
| NR3C2 | HSP90AA1 | 4442077 | 4440643 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.474 | 0.9 | 0.425 | 0.967 |
| AKR1C3 | CBR1 | 4445414 | 4436857 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.063 | 0 | 0.9 | 0.668 | 0.966 |
| ESR1 | HSP90AB1 | 4448358 | 4443770 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.474 | 0.9 | 0.421 | 0.966 |
| MMP9 | MMP8 | 4443919 | 4433667 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0.738 | 0.546 | 0 | 0.9 | 0.915 | 0.964 |
| CRABP2 | RARG | 4451723 | 4447554 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.098 | 0.064 | 0.9 | 0.63 | 0.964 |
| AKT2 | PIK3CG | 4445979 | 4442215 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.063 | 0.174 | 0.9 | 0.594 | 0.964 |
| ESR1 | PGR | 4448358 | 4439740 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.474 | 0.9 | 0.964 | 0.964 |
| ESR1 | CYP19A1 | 4448358 | 4446543 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.085 | 0 | 0.96 | 0.962 |
| CHEK1 | CDK2 | 4447561 | 4435732 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0.308 | 0.719 | 0.325 | 0.261 | 0.9 | 0.885 | 0.962 |
| KAT2B | MDM2 | 4443531 | 4434604 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.379 | 0.9 | 0.426 | 0.961 |
| MMP9 | MMP1 | 4443919 | 4439553 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0.75 | 0.518 | 0 | 0.9 | 0.91 | 0.961 |
| CCNA2 | MDM2 | 4436109 | 4434604 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.062 | 0.108 | 0.9 | 0.58 | 0.96 |
| GSTM1 | GSTM2 | 4438694 | 4433798 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0.447 | 0.981 | 0.83 | 0.379 | 0.65 | 0.831 | 0.96 |
| HSPA8 | EGFR | 4449616 | 4436186 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.393 | 0.9 | 0.386 | 0.959 |
| MMP9 | MMP3 | 4443919 | 4437610 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0.791 | 0.518 | 0 | 0.9 | 0.911 | 0.959 |
| CDK6 | CDK7 | 4435642 | 4434483 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0.393 | 0.846 | 0.063 | 0.504 | 0.9 | 0.805 | 0.958 |
| CTSK | MMP13 | 4435989 | 4434767 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.062 | 0 | 0.9 | 0.591 | 0.958 |
| MMP9 | CTSG | 4443919 | 4432948 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.089 | 0 | 0.9 | 0.569 | 0.957 |
| CDK2 | CDK7 | 4435732 | 4434485 | 9606. ENSP09606. ENSP0 | 0 | 0 | 0.408 | 0.926 | 0.067 | 0.539 | 0.9 | 0.92 | 0.957 |
| RXRA | PPARD | 4449061 | 4438644 | 9606. ENSP09606. ENSP0 | | | | | | | | | |

| | | | | | | | | | | | | |
|----------|---------|---------|--------------------------------|-------|---|-------|-------|-------|-------|-------|-------|-------|
| HSP90AB1 | NOS3 | 4443770 | 4437409 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.06 | 0 | 0.9 | 0.525 | 0.951 |
| AKT2 | MDM2 | 4445979 | 4434604 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.05 | 0.9 | 0.529 | 0.951 | |
| PLAU | EGFR | 4444017 | 4436180 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.088 | 0 | 0.9 | 0.513 | 0.951 |
| ELANE | RNASE2 | 4450799 | 4437950 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.335 | 0 | 0.9 | 0.315 | 0.95 |
| ERBB4 | MDM2 | 4441238 | 4434604 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.305 | 0.9 | 0.33 | 0.949 |
| EPHX2 | CYP2C8 | 4449539 | 4443714 9606. ENSP09606. ENSP0 | 0.056 | 0 | 0 | 0 | 0.051 | 0.081 | 0.9 | 0.481 | 0.949 |
| LCN2 | MMP8 | 4444080 | 4433667 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.097 | 0 | 0.9 | 0.473 | 0.948 |
| ESR1 | KAT2B | 4448358 | 4435311 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.194 | 0.9 | 0.398 | 0.947 |
| ELANE | LCN2 | 4450799 | 4440480 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.12 | 0 | 0.9 | 0.442 | 0.946 |
| HSP90AB1 | PGR | 4443770 | 4439740 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.299 | 0.9 | 0.296 | 0.946 |
| HSP90AA1 | AURKA | 4440643 | 4432988 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.072 | 0.092 | 0.9 | 0.444 | 0.946 |
| LYZ | CTSG | 4434847 | 4432948 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.213 | 0 | 0.9 | 0.378 | 0.946 |
| KAT2B | NR3C1 | 4435311 | 4433539 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.083 | 0.194 | 0.9 | 0.349 | 0.945 |
| PAH | DHFR | 4450247 | 4447922 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.062 | 0 | 0.9 | 0.463 | 0.945 |
| AR | NR3C2 | 4444436 | 4442077 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0.779 | 0.061 | 0.379 | 0.9 | 0.67 | 0.945 |
| AR | NR3C1 | 4444436 | 4433539 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0.816 | 0 | 0.379 | 0.9 | 0.806 | 0.944 |
| HSPA8 | NR3C1 | 4449616 | 4433539 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.166 | 0.9 | 0.386 | 0.944 |
| ESR1 | EGFR | 4448358 | 4436180 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.402 | 0 | 0.908 | 0.943 |
| ELANE | MMP1 | 4450799 | 4439553 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.052 | 0 | 0.9 | 0.45 | 0.943 |
| NR3C2 | NR3C1 | 4442077 | 4433539 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0.861 | 0 | 0.379 | 0.9 | 0.942 | 0.943 |
| RNASE2 | CTSG | 4437950 | 4432948 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.333 | 0 | 0.9 | 0.205 | 0.942 |
| CRABP2 | RXRA | 4451723 | 4449061 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.064 | 0.9 | 0.432 | 0.942 |
| ADK | AHCY | 4436683 | 4433021 9606. ENSP09606. ENSP0 | 0.135 | 0 | 0 | 0 | 0.389 | 0.28 | 0 | 0.863 | 0.941 |
| NR3C1 | PPP5C | 4433539 | 4432591 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.311 | 0.9 | 0.225 | 0.941 |
| CHEK1 | TYMS | 4447561 | 4439011 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.811 | 0.275 | 0 | 0.597 | 0.94 |
| HSP90AB1 | NR3C2 | 4443770 | 4442077 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.311 | 0.9 | 0.201 | 0.94 |
| HSP90AA1 | IL2 | 4440643 | 4433370 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0 | 0.9 | 0.416 | 0.939 |
| PNP | TK1 | 4442485 | 4437779 9606. ENSP09606. ENSP0 | 0.21 | 0 | 0 | 0 | 0.081 | 0 | 0.9 | 0.239 | 0.937 |
| HSPA8 | AR | 4449616 | 4444436 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.109 | 0.9 | 0.348 | 0.936 |
| MMP1 | CTSG | 4439553 | 4432948 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0 | 0.9 | 0.389 | 0.936 |
| MMP9 | CTSS | 4443919 | 4443237 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.134 | 0 | 0.9 | 0.319 | 0.935 |
| LYZ | MMP8 | 4434847 | 4433667 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.116 | 0 | 0.9 | 0.325 | 0.935 |
| NQO1 | ADK | 4439310 | 4436683 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.869 | 0 | 0.527 | 0.935 |
| NMNAT1 | BST1 | 4444851 | 4435534 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0 | 0.9 | 0.377 | 0.935 |
| ERBB4 | EGFR | 4441238 | 4436180 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0.946 | 0 | 0.347 | 0.9 | 0.905 | 0.935 |
| ELANE | LYZ | 4450799 | 4434847 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.112 | 0 | 0.9 | 0.323 | 0.934 |
| GART | TYMS | 4445605 | 4439011 9606. ENSP09606. ENSP0 | 0.291 | 0 | 0 | 0 | 0.14 | 0 | 0 | 0.899 | 0.933 |
| RXRA | KAT2B | 4449061 | 4435311 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.061 | 0.194 | 0.9 | 0.222 | 0.933 |
| RNASE2 | LYZ | 4437950 | 4434847 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.263 | 0 | 0.9 | 0.171 | 0.933 |
| AKT2 | NOS3 | 4445979 | 4437409 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.049 | 0.065 | 0.9 | 0.326 | 0.932 |
| CDK2 | AURKA | 4435732 | 4432988 9606. ENSP09606. ENSP0 | 0 | 0 | 0.431 | 0.76 | 0.303 | 0.88 | 0 | 0.587 | 0.932 |
| LCN2 | LYZ | 4444080 | 4434847 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.101 | 0 | 0.9 | 0.297 | 0.931 |
| HSPA8 | PGR | 4449616 | 4439740 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.183 | 0.9 | 0.229 | 0.931 |
| MMP9 | LYZ | 4443919 | 4434847 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.134 | 0 | 0.9 | 0.251 | 0.929 |
| ESR1 | PPP5C | 4448358 | 4432591 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.324 | 0.9 | 0 | 0.929 |
| DHFR | TPH1 | 4447922 | 4434115 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.061 | 0 | 0.9 | 0.289 | 0.927 |
| HCK | F2 | 4450141 | 4438421 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.05 | 0.9 | 0.287 | 0.926 |
| CHEK1 | CDK7 | 4447561 | 4434485 9606. ENSP09606. ENSP0 | 0 | 0 | 0.31 | 0.642 | 0.062 | 0 | 0.9 | 0.493 | 0.926 |
| ESR1 | NR3C1 | 4448358 | 4433539 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0.66 | 0 | 0 | 0.9 | 0.751 | 0.925 |
| RXRA | ACADM | 4449061 | 4443625 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.061 | 0 | 0.9 | 0.268 | 0.925 |
| PNP | BST1 | 4442485 | 4435534 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0 | 0.9 | 0.271 | 0.924 |
| RXRA | NR3C1 | 4449061 | 4433539 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0.632 | 0 | 0 | 0.9 | 0.641 | 0.923 |
| RXRA | PCK1 | 4449061 | 4439313 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.061 | 0.05 | 0.9 | 0.239 | 0.923 |
| MMP9 | EPHB4 | 4443919 | 4442083 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0 | 0.9 | 0.269 | 0.923 |
| EPHA2 | EPHB4 | 4442119 | 4442085 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0.919 | 0.126 | 0.128 | 0.9 | 0.691 | 0.921 |
| IGF1R | MDM2 | 4435797 | 4434604 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.56 | 0.6 | 0.589 | 0.921 |
| CTSS | MMP8 | 4443237 | 4433667 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.09 | 0 | 0.9 | 0.194 | 0.92 |
| MMP9 | MMP13 | 4443919 | 4434767 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0.797 | 0.064 | 0 | 0.9 | 0.878 | 0.919 |
| AR | PGR | 4444436 | 4439740 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0.813 | 0.069 | 0 | 0.9 | 0.904 | 0.919 |
| LCN2 | MMP9 | 4444080 | 4434919 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.095 | 0.379 | 0 | 0.867 | 0.919 |
| HSPA8 | NR3C2 | 4449616 | 4442077 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.109 | 0.9 | 0.163 | 0.919 |
| AKR1C3 | SULT2B1 | 4445414 | 4432786 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0 | 0.9 | 0.221 | 0.918 |
| RARG | KAT2B | 4447554 | 4435311 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.194 | 0.9 | 0.043 | 0.916 |
| MAPK10 | NR3C1 | 4442217 | 4433539 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.051 | 0 | 0.9 | 0.179 | 0.915 |
| VDR | RARG | 4450226 | 4447554 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0.705 | 0.061 | 0 | 0.9 | 0.466 | 0.914 |
| NR3C2 | PGR | 4442077 | 4439740 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0.809 | 0.061 | 0 | 0.9 | 0.663 | 0.914 |
| CHEK1 | KIT | 4447561 | 4436763 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0.554 | 0 | 0.057 | 0.9 | 0.296 | 0.913 |
| TYMS | KIF11 | 4439011 | 4434801 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.897 | 0 | 0 | 0.181 | 0.912 |
| PGR | NR3C1 | 4439740 | 4433539 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0.839 | 0 | 0 | 0.9 | 0.78 | 0.912 |
| HCK | ITK | 4450141 | 4448039 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0.845 | 0.081 | 0 | 0.9 | 0.501 | 0.911 |
| MMP3 | MMP13 | 4437610 | 4434767 9606. ENSP09606. ENSP0 | 0 | 0 | 0.249 | 0.944 | 0.085 | 0 | 0.9 | 0.893 | 0.91 |
| MMP1 | MMP13 | 4439553 | 4434767 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0.935 | 0.077 | 0 | 0.9 | 0.863 | 0.909 |
| MAPK10 | BCL2L1 | 4442217 | 4437866 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.053 | 0.8 | 0.554 | 0.908 |
| HCK | EPHA2 | 4450141 | 4442119 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0.77 | 0.061 | 0 | 0.9 | 0.255 | 0.907 |
| TYMS | CCNA2 | 4439011 | 4436109 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.838 | 0 | 0 | 0.447 | 0.907 |
| RARG | CDK7 | 4447554 | 4434485 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0 | 0.9 | 0.111 | 0.907 |
| CDA | MMP9 | 4444499 | 4434919 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.089 | 0 | 0.9 | 0.07 | 0.907 |
| RARG | THR8 | 4447554 | 4446583 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0.814 | 0 | 0 | 0.9 | 0.426 | 0.907 |
| PDE4D | ADK | 4441530 | 4436683 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0 | 0.9 | 0.09 | 0.905 |
| PLAU | ITGAL | 4444017 | 4441903 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0 | 0 | 0.9 | 0.089 | 0.905 |
| CDA | LYZ | 4444499 | 4434847 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.095 | 0 | 0.9 | 0 | 0.905 |
| ITGAL | BST1 | 4441903 | 4435534 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.085 | 0 | 0.9 | 0 | 0.904 |
| TYMS | AURKA | 4439011 | 4432988 9606. ENSP09606. ENSP0 | 0 | 0 | 0 | 0 | 0.86 | 0 | 0 | 0.339 | 0.904 |

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|----------|----------|----------|----------|--------|------------|------------|-------|---|-------|-------|-------|-------|-------|-------|-------|-------|
| CHEK1 | HSP90AA1 | 4447561 | 4440643 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0.062 | 0.416 | 0 | 0.803 | 0.883 | |
| AKT2 | BCL2L1 | 4445979 | 4437866 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0.05 | 0 | 0.8 | 0.429 | 0.882 | |
| CHEK1 | KIP11 | 4447561 | 4434801 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0.821 | 0 | 0 | 0.359 | 0.88 | |
| HCK | WAS | 4450141 | 4444767 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0.314 | 0.472 | 0 | 0.694 | 0.879 | |
| CYP19A1 | AR | 4446543 | 4444436 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.085 | 0 | 0.873 | 0.879 | |
| CTSS | CTSB | 4443237 | 4441541 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0.223 | 0.675 | 0.103 | 0 | 0.8 | 0.77 | 0.868 | |
| PGF | TEK | 4450339 | 4445310 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.058 | 0.6 | 0.673 | 0.866 | |
| EGFR | MDM2 | 4436188 | 4434604 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0 | 0 | 0.6 | 0.671 | 0.862 | |
| BCL2L1 | CDK2 | 4437866 | 4435732 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0.05 | 0.13 | 0 | 0.845 | 0.861 | |
| TNK2 | AR | 4445619 | 4444436 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.379 | 0 | 0.781 | 0.858 | |
| AKT2 | HSP90AB1 | 4445979 | 4443770 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0.061 | 0.584 | 0.6 | 0.201 | 0.858 | |
| CYP19A1 | PGR | 4446543 | 4439740 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.085 | 0 | 0.851 | 0.857 | |
| DHFR | DHDH | 4447922 | 4433071 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0.08 | 0 | 0 | 0.849 | 0.855 | |
| ESR1 | MAPK10 | 4448358 | 4442217 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0 | 0 | 0.8 | 0.293 | 0.852 | |
| SDS | AHCY | 4434551 | 4433021 | 9606. | ENSP09606. | ENSP0 | 0.043 | 0 | 0 | 0 | 0.116 | 0 | 0.8 | 0.221 | 0.85 | |
| ADAM17 | EGFR | 4438549 | 4436186 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0.061 | 0 | 0 | 0.842 | 0.846 | |
| CHEK1 | AURKA | 4447561 | 4432988 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0.288 | 0.675 | 0.754 | 0.162 | 0 | 0.672 | 0.845 | |
| BCL2L1 | ANXA5 | 4437866 | 4437297 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0.837 | 0.837 | |
| GLO1 | SDS | 4441469 | 4434551 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.055 | 0.8 | 0.203 | 0.836 | |
| GART | DHDH | 4445605 | 4433071 | 9606. | ENSP09606. | ENSP0 | 0.37 | 0 | 0 | 0 | 0.311 | 0.057 | 0 | 0.644 | 0.835 | |
| GART | IMPDH1 | 4445605 | 4441488 | 9606. | ENSP09606. | ENSP0 | 0.353 | 0 | 0 | 0 | 0.261 | 0 | 0 | 0.683 | 0.835 | |
| GART | APRT | 4445605 | 4445041 | 9606. | ENSP09606. | ENSP0 | 0.367 | 0 | 0 | 0 | 0.082 | 0.076 | 0 | 0.724 | 0.832 | |
| MMP9 | EGFR | 4443919 | 4436186 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0.828 | 0.828 | |
| TK1 | CCNA2 | 4437779 | 4436109 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0.698 | 0 | 0 | 0.454 | 0.828 | |
| TYMS | CDK2 | 4439011 | 4435732 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0.686 | 0 | 0 | 0.467 | 0.825 | |
| MAPK10 | PRKCQ | 4442217 | 4435203 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0.588 | 0 | 0.157 | 0.8 | 0.048 | 0.824 | |
| HSP90AB1 | EGFR | 4443770 | 4436186 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.487 | 0 | 0.668 | 0.823 | |
| MAOB | GLO1 | 4444997 | 4441169 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0 | 0 | 0.8 | 0.133 | 0.819 | |
| MAPK10 | PGR | 4442217 | 4439740 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0.08 | 0 | 0.8 | 0.082 | 0.816 | |
| PDE5A | NOS3 | 4441685 | 4437409 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0.814 | 0.814 | |
| PGR | EGFR | 4439740 | 4436186 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.077 | 0 | 0.804 | 0.811 | |
| EGFR | IL2 | 4436186 | 4433376 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0 | 0 | 0.6 | 0.548 | 0.811 | |
| CDK2 | KIP11 | 4435732 | 4434801 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0.373 | 0.162 | 0 | 0.666 | 0.809 | |
| ELANE | MMP9 | 4450799 | 4443919 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0.094 | 0 | 0 | 0.794 | 0.805 | |
| MTAP | APRT | 4445335 | 4445041 | 9606. | ENSP09606. | ENSP0 | 0.219 | 0 | 0 | 0 | 0.064 | 0 | 0 | 0.753 | 0.803 | |
| AKR1C2 | MIF | 4445450 | 4432873 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.794 | 0 | 0.086 | 0.803 | |
| AR | EGFR | 4444436 | 4436186 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0.062 | 0.402 | 0 | 0.678 | 0.803 | |
| EGFR | LGALS3 | 4436186 | 4434352 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0.076 | 0.379 | 0 | 0.685 | 0.803 | |
| CASP7 | BCL2L1 | 4443299 | 4437866 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.168 | 0 | 0.765 | 0.796 | |
| ADH1C | GSTM1 | 4449347 | 4438694 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0.061 | 0 | 0.65 | 0.428 | 0.795 | |
| GSTM1 | ADH1B | 4438694 | 4438237 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0.061 | 0 | 0.65 | 0.413 | 0.79 | |
| MMP9 | CTSK | 4443919 | 4435983 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0.089 | 0 | 0 | 0.779 | 0.79 | |
| CASP7 | BIRC7 | 4443299 | 4433001 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0.048 | 0.57 | 0 | 0.528 | 0.79 | |
| GSTP1 | CYP2C8 | 4446837 | 443714 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.077 | 0.65 | 0.399 | 0.789 | |
| PGF | KIT | 4450339 | 4436763 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.058 | 0.6 | 0.481 | 0.787 | |
| PPARG | EGFR | 4436745 | 4436186 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0.062 | 0.402 | 0 | 0.65 | 0.787 | |
| CYP2C8 | GSTM1 | 4443714 | 4438694 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.077 | 0.65 | 0.391 | 0.786 | |
| PIK3CG | KIT | 4442215 | 4436763 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.719 | 0 | 0.267 | 0.785 | |
| AR | IGF1R | 4444436 | 4435797 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0.051 | 0.085 | 0 | 0.77 | 0.783 | |
| MMP9 | NOS3 | 4443919 | 4437409 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0.062 | 0 | 0 | 0.778 | 0.783 | |
| PGF | EGFR | 4450339 | 4436186 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0 | 0 | 0.6 | 0.473 | 0.78 | |
| ESR1 | AKT2 | 4448358 | 4445979 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.407 | 0 | 0.638 | 0.777 | |
| CASP1 | BCL2L1 | 4449658 | 4437866 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.461 | 0 | 0.596 | 0.773 | |
| AKR1C3 | AR | 4445414 | 4444436 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.05 | 0 | 0.769 | 0.772 | |
| APRT | TK1 | 4445041 | 4437779 | 9606. | ENSP09606. | ENSP0 | 0.174 | 0 | 0 | 0 | 0.136 | 0 | 0 | 0.699 | 0.767 | |
| APRT | CDA | 4445041 | 4444499 | 9606. | ENSP09606. | ENSP0 | 0.23 | 0 | 0 | 0 | 0 | 0 | 0 | 0.704 | 0.762 | |
| NOS3 | PPARG | 4437409 | 4436745 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0.053 | 0.091 | 0 | 0.741 | 0.757 | |
| DHFR | TK1 | 4447922 | 4437779 | 9606. | ENSP09606. | ENSP0 | 0.133 | 0 | 0 | 0 | 0.227 | 0 | 0 | 0.664 | 0.756 | |
| GART | CDA | 4445605 | 4444498 | 9606. | ENSP09606. | ENSP0 | 0.326 | 0 | 0 | 0 | 0 | 0 | 0 | 0.653 | 0.756 | |
| ESR1 | ERBB4 | 4448358 | 4441238 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.402 | 0 | 0.605 | 0.754 | |
| MMP9 | LGALS3 | 4443919 | 4434352 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0.071 | 0 | 0 | 0.743 | 0.751 | |
| AR | MET | 4444436 | 4439114 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.085 | 0 | 0.738 | 0.75 | |
| CTSB | BCL2L1 | 4441541 | 4437866 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0.044 | 0 | 0.6 | 0.398 | 0.749 | |
| PGF | MMP9 | 4450339 | 4446837 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.65 | 0.65 | 0.291 | 0.741 | |
| ESR1 | ANXA5 | 4443919 | 4441169 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.055 | 0 | 0.364 | 0.733 | |
| TYMS | EGFR | 44437297 | 4436186 | 9606. | ENSP09606. | ENSP0 | 0.07 | 0 | 0 | 0 | 0.064 | 0.413 | 0 | 0.548 | 0.738 | |
| BMP2 | MMP13 | 4445120 | 4444767 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0.736 | 0.736 | |
| HSP90AA1 | CDK2 | 4440643 | 4435732 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0.054 | 0.414 | 0 | 0.561 | 0.735 | |
| ADH1C | MAOB | 4449347 | 4444997 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0.092 | 0 | 0.65 | 0.23 | 0.734 | |
| PGF | MET | 4450339 | 44450339 | 947790 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.058 | 0.6 | 0.348 | 0.732 |
| MMP1 | LGALS3 | 4439553 | 4434352 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0.097 | 0 | 0 | 0.715 | 0.732 | |
| CASP7 | ANXA5 | 4443299 | 4437297 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0.732 | 0.732 | |
| BCL2L1 | MDM2 | 4437866 | 4434604 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0.055 | 0.189 | 0 | 0.675 | 0.729 | |
| TK1 | AURKA | 4437779 | 4432988 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0.618 | 0 | 0 | 0.31 | 0.725 | |
| PLAU | MMP9 | 4444017 | 4443919 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0.076 | 0 | 0 | 0.713 | 0.724 | |
| CDA | TYMS | 4444499 | 4439011 | 9606. | ENSP09606. | ENSP0 | 0.052 | 0 | 0 | 0 | 0.061 | 0 | 0 | 0.713 | 0.723 | |
| GSTP1 | NQ01 | 4446837 | 4439310 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0.064 | 0 | 0 | 0.717 | 0.723 | |
| ESR1 | MDM2 | 4448358 | 4434604 | 9606. | ENSP09606. | ENSP0</td | | | | | | | | | | |

| | | | | | | | | | | | | | | | |
|----------|----------|---------|---------|-------|------------|-------|-------|---|---|-------|-------|-------|------|-------|-------|
| THR8 | KAT2B | 4446585 | 4435311 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0.061 | 0.194 | 0.6 | 0.177 | 0.717 |
| MET | EGFR | 4439114 | 4436186 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0.58 | 0.229 | 0.418 | 0 | 0.933 | 0.715 |
| AKT2 | CCNA2 | 4445979 | 4436109 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.354 | 0 | 0.571 | 0.711 |
| CHEK1 | HSP90AB1 | 4447561 | 4443770 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0.062 | 0.416 | 0 | 0.513 | 0.71 |
| MAOB | BCHE | 4444997 | 4435424 | 9606. | ENSP09606. | ENSP0 | 0.045 | 0 | 0 | 0 | 0.086 | 0 | 0 | 0.694 | 0.71 |
| CYP2C8 | GSTM2 | 4443714 | 4433798 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.077 | 0.65 | 0.176 | 0.71 |
| MMP9 | PPARG | 4443919 | 4436745 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0.709 | 0.709 |
| MAOB | ADH1B | 4444997 | 4438237 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0.104 | 0 | 0.65 | 0.142 | 0.707 |
| EPHA2 | HSP90AA1 | 4442119 | 4440643 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0.048 | 0.244 | 0 | 0.625 | 0.707 |
| HSP90AA1 | MDM2 | 4440643 | 4434604 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.348 | 0 | 0.568 | 0.706 |
| AR | BCL2L1 | 4444436 | 4437866 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.064 | 0 | 0.698 | 0.706 |
| GSTP1 | GSTM2 | 4446837 | 4433798 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0.803 | 0 | 0 | 0.65 | 0.801 | 0.704 |
| HSP90AA1 | MET | 4440643 | 4439114 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0 | 0.452 | 0 | 0.483 | 0.704 |
| PGF | IGF1R | 4450339 | 4435797 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0 | 0 | 0.6 | 0.287 | 0.702 |
| F2 | BCHE | 4438421 | 4435424 | 9606. | ENSP09606. | ENSP0 | 0 | 0 | 0 | 0 | 0.126 | 0.052 | 0 | 0.669 | 0.701 |

Table 6S

Top 15 in network 0.7PPIstring_interactions ranked by degree method

| Rank | Name | Score |
|------|----------|-------|
| 1 | EGFR | 20 |
| 2 | MMP9 | 19 |
| 3 | HSP90AA1 | 18 |
| 4 | AR | 17 |
| 5 | ESR1 | 15 |
| 6 | MDM2 | 13 |
| 7 | HSP90AB1 | 12 |
| 8 | TYMS | 11 |
| 8 | CDK2 | 11 |
| 8 | RXRA | 11 |
| 8 | PGR | 11 |
| 8 | NR3C1 | 11 |
| 13 | CCNA2 | 10 |
| 13 | CHEK1 | 10 |
| 15 | BCL2L1 | 9 |

Table 7S

| Pw | Count | PValue | FDR |
|------------|-------|-----------|-----------|
| cytosol | 77 | 4.21E-20 | 5.30E-17 |
| extracellu | 58 | 6.52E-12 | 8.22E-09 |
| extracellu | 40 | 3.12E-10 | 3.93E-07 |
| extracellu | 35 | 2.02E-09 | 2.54E-06 |
| cytoplasm | 68 | 1.19E-05 | 0.0149911 |
| nucleoplas | 44 | 1.40E-05 | 0.0175977 |
| cell surfa | 16 | 3.60E-05 | 0.0453097 |
| endolysosc | 3 | 3.89E-04 | 0.4890617 |
| receptor c | 7 | 5.98E-04 | 0.7513211 |
| extracellu | 10 | 7.13E-04 | 0.8944314 |
| focal adhe | 10 | 0.00471 | 5.7772062 |
| mitochondr | 9 | 0.0053124 | 6.4934207 |
| membrane | 30 | 0.005562 | 6.788777 |
| myelin she | 6 | 0.0080481 | 9.6837811 |
| melanosome | 5 | 0.0092819 | 11.089567 |
| cell-cell | 6 | 0.0132426 | 15.467336 |
| intracellu | 11 | 0.015806 | 18.193978 |
| basolatera | 6 | 0.0158392 | 18.228727 |
| cyclin A2- | 2 | 0.0161768 | 18.581572 |
| nucleus | 57 | 0.0203152 | 22.794604 |
| nuclear ch | 6 | 0.0207427 | 23.218127 |
| mitochondr | 19 | 0.0226223 | 25.055208 |
| ruffle mem | 4 | 0.0291687 | 31.142207 |
| RNA polyme | 3 | 0.0293451 | 31.299724 |
| apical pla | 7 | 0.0316746 | 33.34926 |
| lysosomal | 4 | 0.0319701 | 33.605116 |
| lysosome | 6 | 0.0373457 | 38.105618 |
| spindle mi | 3 | 0.0496366 | 47.360724 |
| leading ed | 2 | 0.0631641 | 56.062857 |
| proteinace | 6 | 0.0677539 | 58.700332 |

Table 8S

| Pw | Count | PValue | FDR |
|-------------|-------|-----------|-----------|
| steroid hc | 13 | 5.71E-14 | 8.15E-11 |
| serine-typ | 17 | 9.39E-10 | 1.34E-06 |
| transmembr | 9 | 1.19E-09 | 1.70E-06 |
| ATP bindin | 38 | 5.48E-09 | 7.83E-06 |
| protein ty | 12 | 2.62E-08 | 3.75E-05 |
| zinc ion b | 32 | 2.65E-08 | 3.79E-05 |
| identical | 24 | 1.71E-07 | 2.44E-04 |
| drug bindi | 9 | 3.46E-07 | 4.95E-04 |
| RNA polym | 7 | 6.69E-07 | 9.56E-04 |
| endopeptid | 7 | 7.79E-06 | 0.0111211 |
| enzyme bin | 14 | 8.15E-06 | 0.0116465 |
| protein ki | 14 | 1.81E-05 | 0.025886 |
| oxidoreduc | 4 | 3.65E-05 | 0.0521733 |
| nitric-oxi | 4 | 3.65E-05 | 0.0521733 |
| steroid bi | 5 | 8.73E-05 | 0.1246917 |
| cysteine-t | 6 | 2.00E-04 | 0.2855515 |
| phosphatid | 6 | 2.16E-04 | 0.3083788 |
| ketosteroi | 3 | 2.31E-04 | 0.3291885 |
| retinoid X | 4 | 2.84E-04 | 0.4044211 |
| cysteine-t | 5 | 3.41E-04 | 0.485572 |
| oxidoreduc | 9 | 4.04E-04 | 0.5756161 |
| phenanthre | 3 | 4.59E-04 | 0.6535665 |
| trans-1, 2- | 3 | 4.59E-04 | 0.6535665 |
| metalloend | 7 | 4.97E-04 | 0.7079339 |
| Ras guanyl | 7 | 5.46E-04 | 0.7768751 |
| protein hc | 17 | 7.32E-04 | 1.0401524 |
| kinase act | 9 | 0.0013605 | 1.9258602 |
| alditol:NA | 3 | 0.0015786 | 2.2313228 |
| carboxylic | 3 | 0.0015786 | 2.2313228 |
| collagen b | 5 | 0.0019372 | 2.7317817 |
| bile acid | 3 | 0.0020926 | 2.9478305 |
| sequence-s | 13 | 0.0021673 | 3.0515356 |
| protein bi | 96 | 0.0024633 | 3.4614612 |
| virion bin | 3 | 0.0033244 | 4.6449806 |
| receptor s | 3 | 0.0033244 | 4.6449806 |
| glutathion | 3 | 0.0040396 | 5.6178193 |
| proteoglyc | 3 | 0.0040396 | 5.6178193 |
| receptor b | 10 | 0.0041547 | 5.7735066 |
| catalytic | 7 | 0.0065051 | 8.9007705 |
| MHC class | 3 | 0.0085628 | 11.558993 |
| scaffold p | 4 | 0.0087198 | 11.758923 |
| transferas | 5 | 0.0103444 | 13.802179 |
| retinol de | 3 | 0.0107924 | 14.357869 |
| iron ion b | 6 | 0.0115478 | 15.287367 |
| monooxygen | 4 | 0.014595 | 18.942125 |
| CTP bindin | 2 | 0.0175756 | 22.374699 |
| protein ph | 4 | 0.0181958 | 23.071764 |
| protein ki | 9 | 0.0185702 | 23.489676 |
| protein se | 9 | 0.0185702 | 23.489676 |

| | | | |
|--------------|---|-----------|-----------|
| glycoprote | 4 | 0.019762 | 24.806138 |
| indanol de | 2 | 0.026248 | 31.608829 |
| ICAM-3 rec | 2 | 0.026248 | 31.608829 |
| UTP bindin | 2 | 0.026248 | 31.608829 |
| sulfonylur | 2 | 0.026248 | 31.608829 |
| vitamin D | 2 | 0.026248 | 31.608829 |
| ATPase bin | 4 | 0.0277036 | 33.054738 |
| carbohydr | 6 | 0.0300332 | 35.309726 |
| nucleoside | 2 | 0.0348443 | 39.744942 |
| dATP bindi | 2 | 0.0348443 | 39.744942 |
| Rho GDP-di | 2 | 0.0348443 | 39.744942 |
| metallopep | 4 | 0.0348878 | 39.783707 |
| cyclin-dep | 3 | 0.036107 | 40.861167 |
| glutathion | 3 | 0.0380779 | 42.565169 |
| ubiquitin | 7 | 0.0420763 | 45.882689 |
| TPR domain | 2 | 0.0433653 | 46.913548 |
| fibroblast | 2 | 0.0433653 | 46.913548 |
| arachidoni | 2 | 0.0433653 | 46.913548 |
| retinoic a | 2 | 0.0433653 | 46.913548 |
| electron c | 4 | 0.0453963 | 48.500901 |
| peptidase | 4 | 0.0453963 | 48.500901 |
| alcohol de | 2 | 0.0518115 | 53.22965 |
| arylesterase | 2 | 0.0518115 | 53.22965 |
| heat shock | 3 | 0.0529549 | 54.028843 |
| heparin bi | 5 | 0.0534475 | 54.3692 |
| growth fac | 5 | 0.0550198 | 55.439985 |
| 1-phosphat | 3 | 0.0552247 | 55.577812 |
| small mole | 2 | 0.0601837 | 58.794588 |
| non-membra | 3 | 0.0622336 | 60.05996 |
| retinoid b | 2 | 0.0684824 | 63.697659 |
| thyroid hc | 2 | 0.0684824 | 63.697659 |
| unfolded p | 4 | 0.0736087 | 66.449238 |
| ion channe | 4 | 0.0783823 | 68.835896 |
| retinal bi | 2 | 0.0848621 | 71.823597 |
| ephrin rec | 2 | 0.0929443 | 75.176896 |
| peptide bi | 3 | 0.0956377 | 76.209206 |

Table 9S

| Pw | Count | PValue | FDR |
|------------|-------|-----------|-----------|
| steroid hc | 11 | 6.31E-11 | 1.06E-07 |
| negative r | 20 | 1.95E-08 | 3.28E-05 |
| positive r | 18 | 7.70E-07 | 0.0012942 |
| positive r | 7 | 1.30E-05 | 0.0218364 |
| response t | 13 | 1.62E-05 | 0.0272914 |
| positive r | 8 | 1.33E-04 | 0.2226868 |
| glucose me | 6 | 3.09E-04 | 0.518887 |
| positive r | 10 | 5.31E-04 | 0.8898674 |
| positive r | 6 | 8.77E-04 | 1.4649836 |
| response t | 6 | 9.26E-04 | 1.5450522 |
| positive r | 5 | 0.0013016 | 2.1663835 |
| endocytosi | 7 | 0.0014537 | 2.4167745 |
| innate imm | 12 | 0.0015107 | 2.5102745 |
| positive r | 4 | 0.0020723 | 3.4283755 |
| daunorubic | 3 | 0.0020866 | 3.4516013 |
| doxorubici | 3 | 0.0020866 | 3.4516013 |
| mesenchyma | 3 | 0.0020866 | 3.4516013 |
| angiogenes | 8 | 0.0036501 | 5.9642369 |
| regulation | 3 | 0.0040281 | 6.5623449 |
| response t | 7 | 0.0042126 | 6.8530181 |
| positive r | 7 | 0.0045768 | 7.4242074 |
| cell proli | 10 | 0.0051941 | 8.3850301 |
| positive r | 5 | 0.0057026 | 9.1692703 |
| positive r | 7 | 0.0058311 | 9.3664356 |
| negative r | 14 | 0.0112413 | 17.313224 |
| nucleoside | 3 | 0.0119594 | 18.31734 |
| G1/S trans | 5 | 0.0126462 | 19.26696 |
| negative r | 4 | 0.0132274 | 20.062381 |
| cellular c | 3 | 0.0145192 | 21.804169 |
| regulation | 4 | 0.0181254 | 26.479321 |
| Notch sign | 5 | 0.0188582 | 27.396706 |
| leukocyte | 3 | 0.0202717 | 29.135693 |
| cellular p | 5 | 0.0205175 | 29.434061 |
| drug metab | 3 | 0.0234481 | 32.901895 |
| regulation | 4 | 0.0239022 | 33.424563 |
| negative r | 9 | 0.0242139 | 33.781283 |
| regulation | 6 | 0.0243397 | 33.924619 |
| regulation | 3 | 0.025108 | 34.794227 |
| Fc-gamma r | 5 | 0.0260224 | 35.815149 |
| apoptotic | 11 | 0.0286708 | 38.687859 |
| G2/M trans | 5 | 0.0330569 | 43.181063 |
| cell surfa | 7 | 0.034598 | 44.684902 |
| regulation | 2 | 0.0347946 | 44.873992 |
| cell divis | 8 | 0.0356054 | 45.647567 |
| extrinsic | 3 | 0.036011 | 46.030749 |
| positive r | 10 | 0.0388349 | 48.628704 |
| deoxyribon | 2 | 0.0433037 | 52.500831 |
| mitotic ce | 2 | 0.0433037 | 52.500831 |
| negative r | 3 | 0.0441129 | 53.17199 |

| | | | |
|------------|---|-----------|------------|
| negative r | 3 | 0.0483932 | 56. 576099 |
| pyrimidine | 2 | 0.0517382 | 59. 07284 |
| tetrahydrc | 2 | 0.0517382 | 59. 07284 |
| positive r | 3 | 0.0528178 | 59. 849383 |
| cellular r | 4 | 0.0529858 | 59. 968989 |
| T cell act | 3 | 0.0644692 | 67. 394155 |
| humoral ir | 3 | 0.0899906 | 79. 521824 |
| regulation | 4 | 0.0966711 | 81. 908317 |

Table 10S

| Pw | Count | PValue | FDR |
|-------------|-------|-----------|-----------|
| Pathways i | 26 | 1.82E-08 | 2.30E-05 |
| PI3K-Akt s | 21 | 2.57E-06 | 0.0032423 |
| Prostate c | 11 | 3.22E-06 | 0.0040628 |
| Progester c | 9 | 1.49E-04 | 0.188181 |
| Ras signal | 14 | 1.84E-04 | 0.2315246 |
| Melanoma | 8 | 2.54E-04 | 0.3198339 |
| Metabolism | 8 | 3.28E-04 | 0.4138799 |
| Rap1 signa | 13 | 3.49E-04 | 0.4396727 |
| Estrogen s | 9 | 3.66E-04 | 0.4605876 |
| Metabolic | 38 | 4.98E-04 | 0.6265678 |
| Proteoglyc | 12 | 8.53E-04 | 1.0713798 |
| Central ca | 7 | 9.30E-04 | 1.1671398 |
| Pancreatic | 7 | 0.0010093 | 1.2667145 |
| Drug metab | 7 | 0.0012805 | 1.6045253 |
| Measles | 9 | 0.0025263 | 3.1428553 |
| Transcript | 10 | 0.0029123 | 3.6149281 |
| Chemical c | 7 | 0.0029547 | 3.6667422 |
| Small cell | 7 | 0.0040007 | 4.9347696 |
| One carbon | 4 | 0.0050922 | 6.2416961 |
| Glioma | 6 | 0.0059139 | 7.2144883 |
| PPAR signa | 6 | 0.0067222 | 8.1624407 |
| Adherens j | 6 | 0.0085656 | 10.291073 |
| FoxO signa | 8 | 0.0099392 | 11.847529 |
| Thyroid hc | 7 | 0.0167224 | 19.17578 |
| Non-small | 5 | 0.0174847 | 19.963265 |
| Steroid hc | 5 | 0.0196568 | 22.168547 |
| Epstein-Ba | 7 | 0.0217359 | 24.226829 |
| Tyrosine n | 4 | 0.0240853 | 26.492365 |
| HIF-1 sign | 6 | 0.0283291 | 30.427264 |
| Viral carc | 9 | 0.0301066 | 32.016962 |
| Epithelial | 5 | 0.0314446 | 33.191488 |
| T cell rec | 6 | 0.0330034 | 34.536245 |
| Pyrimidine | 6 | 0.0342428 | 35.58764 |
| Adipocytok | 5 | 0.0361261 | 37.155535 |
| Phenylalan | 3 | 0.0361997 | 37.216102 |
| Bladder ca | 4 | 0.0363284 | 37.321809 |
| Purine met | 8 | 0.0378885 | 38.590841 |
| Chronic my | 5 | 0.0394595 | 39.844755 |
| TNF signal | 6 | 0.0422851 | 42.04094 |
| Insulin re | 6 | 0.0437275 | 43.133321 |
| Hepatitis | 7 | 0.0450725 | 44.134832 |
| Antigen pr | 5 | 0.0466373 | 45.279609 |
| Fc gamma R | 5 | 0.063027 | 56.038216 |
| AMPK signa | 6 | 0.0689064 | 59.396499 |
| ErbB signa | 5 | 0.0698614 | 59.919124 |
| Cell cycle | 6 | 0.0708211 | 60.438075 |
| Rheumatoid | 5 | 0.0722214 | 61.184175 |
| Focal adhe | 8 | 0.0754628 | 62.861844 |
| NOD-like r | 4 | 0.0781099 | 64.182102 |

| | | | |
|------------|---|-----------|-----------|
| Acute myel | 4 | 0.0781099 | 64.182102 |
| Regulation | 8 | 0.0817074 | 65.90713 |
| Biosynthes | 8 | 0.0849394 | 67.391343 |
| Nicotinate | 3 | 0.0941881 | 71.316212 |
| VEGF signa | 4 | 0.0952433 | 71.735194 |
| Arachidoni | 4 | 0.0952433 | 71.735194 |
| Apoptosis | 4 | 0.0988413 | 73.121837 |