

| id_cp | interacting_partner_a | partner_b | gene_a | gene_b | secreted | receptor_a | receptor_b |
|------------|-----------------------|--------------------------|-----------------|-----------|----------|------------|------------|
| CPI-SC0FA2 | CSPG4_inte | simple:Q6U complex:int | ENSG00000173546 | | FALSE | FALSE | FALSE |
| CPI-SS006C | MDK_LRP1 | simple:P21: simple:Q07 | ENSG00000 | ENSG00000 | TRUE | FALSE | TRUE |
| CPI-SS0B38 | CADM3_EP | simple:Q8N simple:Q9H | ENSG00000 | ENSG00000 | FALSE | FALSE | FALSE |
| CPI-SS0855 | NRP1_PGF | simple:O14 simple:P49 | ENSG00000 | ENSG00000 | TRUE | TRUE | FALSE |
| CPI-CS0E0B | FLT1_comp | complex:FL simple:P49763 | ENSG00000 | | TRUE | TRUE | FALSE |
| CPI-CS0E54 | BMPR1A_B | complex:BN simple:P12644 | ENSG00000 | | TRUE | TRUE | FALSE |
| CPI-SS01F6 | FGF2_FGFR | simple:P09(simple:P11: | ENSG00000 | ENSG00000 | TRUE | FALSE | TRUE |
| CPI-CS0C57 | PlexinA1_cc | complex:Pl simple:Q8NFY4 | ENSG00000 | | FALSE | TRUE | FALSE |
| CPI-SS0CB8 | WNT5A_PT | simple:P41: simple:Q15 | ENSG00000 | ENSG00000 | TRUE | FALSE | TRUE |
| CPI-SS0F21 | SCGB3A1_N | simple:Q96 simple:Q9U | ENSG00000 | ENSG00000 | TRUE | FALSE | TRUE |
| CPI-SC00EA | COL5A3_int | simple:P25: complex:int | ENSG00000080573 | | TRUE | FALSE | FALSE |
| CPI-SC0376 | COL8A1_int | simple:P27(complex:int | ENSG00000144810 | | TRUE | FALSE | FALSE |
| CPI-SC042B | COL4A5_int | simple:P29: complex:int | ENSG00000188153 | | TRUE | FALSE | FALSE |
| CPI-SC0036 | COL3A1_int | simple:P02: complex:int | ENSG00000168542 | | TRUE | FALSE | FALSE |
| CPI-SC0886 | COL4A3_int | simple:Q01 complex:int | ENSG00000169031 | | TRUE | FALSE | FALSE |
| CPI-SC09DB | COL1A2_int | simple:P08: complex:int | ENSG00000164692 | | TRUE | FALSE | FALSE |
| CPI-SC0FB8 | HTR1B_Ser | simple:P28: complex:Se | ENSG00000135312 | | TRUE | TRUE | FALSE |
| CPI-SS068C | FGF1_TGFB | simple:P05: simple:Q03 | ENSG00000 | ENSG00000 | TRUE | FALSE | TRUE |
| CPI-CS0475 | ACVR_1A2A | complex:AC simple:P22003 | ENSG00000 | | TRUE | TRUE | FALSE |
| CPI-CS0FA9 | ACVR_1A2E | complex:AC simple:P22003 | ENSG00000 | | TRUE | TRUE | FALSE |
| CPI-CS047E | BMR1A_AC | complex:BN simple:P22003 | ENSG00000 | | TRUE | TRUE | FALSE |
| CPI-CS0CA3 | BMR1A_AV | complex:BN simple:P22003 | ENSG00000 | | TRUE | TRUE | FALSE |
| CPI-SS0D22 | NRP2_PGF | simple:O60 simple:P49 | ENSG00000 | ENSG00000 | TRUE | TRUE | FALSE |
| CPI-SS033E | NRP1_VEGF | simple:O14 simple:P49 | ENSG00000 | ENSG00000 | TRUE | TRUE | FALSE |
| CPI-SS0DD9 | NGF_SORT1 | simple:P01: simple:Q99 | ENSG00000 | ENSG00000 | TRUE | FALSE | TRUE |
| CPI-SC0B25 | VCAM1_int | simple:P19: complex:int | ENSG00000162692 | | FALSE | FALSE | FALSE |
| CPI-SS02DA | BSG_PPIA | simple:P35(simple:P62: | ENSG00000 | ENSG00000 | TRUE | FALSE | FALSE |
| CPI-SC04AB | COL4A4_int | simple:P53: complex:int | ENSG00000081052 | | TRUE | FALSE | FALSE |
| CPI-CS046B | BMPR1A_B | complex:BN simple:P22003 | ENSG00000 | | TRUE | TRUE | FALSE |
| CPI-SS0A07 | TNFRSF21_ | simple:O75 simple:P05(| ENSG00000 | ENSG00000 | FALSE | TRUE | FALSE |
| CPI-SS0BFB | PLXNB2_PT | simple:O15 simple:P21: | ENSG00000 | ENSG00000 | TRUE | TRUE | FALSE |
| CPI-SS0755 | FGF1_FGFR | simple:P05: simple:P11: | ENSG00000 | ENSG00000 | TRUE | FALSE | TRUE |
| CPI-SS0E61 | FLT1_PGF | simple:P17: simple:P49 | ENSG00000 | ENSG00000 | TRUE | TRUE | FALSE |
| CPI-SS0158 | FGF1_FGFR | simple:P05: simple:P22(| ENSG00000 | ENSG00000 | TRUE | FALSE | TRUE |
| CPI-CS0C01 | FLT1_comp | complex:FL simple:P49765 | ENSG00000 | | TRUE | TRUE | FALSE |
| CPI-SS0EE0 | CDH2_CDH: | simple:P19(simple:P19(| ENSG00000 | ENSG00000 | FALSE | FALSE | FALSE |
| CPI-CS0E1B | ACVR1_BM | complex:AC simple:P22003 | ENSG00000 | | TRUE | TRUE | FALSE |
| CPI-SC03FF | LAMC1_int | simple:P11(complex:int | ENSG00000135862 | | TRUE | FALSE | FALSE |
| CPI-SS0397 | FGFR1_NCA | simple:P11: simple:P13: | ENSG00000 | ENSG00000 | TRUE | TRUE | TRUE |
| CPI-SC0C21 | JAM2_integ | simple:P57(complex:int | ENSG00000154721 | | FALSE | FALSE | FALSE |
| CPI-SS0D95 | PDGFRB_PT | simple:P09(simple:Q9G | ENSG00000 | ENSG00000 | TRUE | TRUE | FALSE |
| CPI-SC0717 | VTN_integr | simple:P04(complex:int | ENSG00000109072 | | TRUE | FALSE | FALSE |
| CPI-SS0F9D | JAM3_JAM: | simple:Q9B simple:Q9B | ENSG00000 | ENSG00000 | TRUE | FALSE | FALSE |

| | | | |
|---|-------|-------|-------|
| CPI-SS087B JAM2_JAM: simple:P57: simple:Q9B ENSG000000 ENSG000000 | TRUE | FALSE | FALSE |
| CPI-SS03A4 AXL_GAS6 simple:P30: simple:Q14 ENSG000000 ENSG000000 | TRUE | TRUE | FALSE |
| CPI-SS0954 CCRL2_RAR simple:O00: simple:Q99 ENSG000000 ENSG000000 | TRUE | TRUE | FALSE |
| CPI-SC09B4 FN1_integri simple:P02: complex:int ENSG000000115414 | TRUE | FALSE | FALSE |
| CPI-SS0BAB ANGPT2_TE simple:O15 simple:Q02 ENSG000000 ENSG000000 | TRUE | FALSE | TRUE |
| CPI-SC0F96 FN1_integri simple:P02: complex:int ENSG000000115414 | TRUE | FALSE | FALSE |
| CPI-SS0415 MERTK_GA: simple:Q12 simple:Q14 ENSG000000 ENSG000000 | TRUE | TRUE | FALSE |
| CPI-SS05A2 PDGFB_PD: simple:P01: simple:P09: ENSG000000 ENSG000000 | TRUE | FALSE | TRUE |
| CPI-SS03E2 ADORA2A_ simple:P29: simple:P49: ENSG000000 ENSG000000 | FALSE | TRUE | FALSE |
| CPI-SS0241 NRP2_SEM: simple:O60: simple:Q99 ENSG000000 ENSG000000 | TRUE | TRUE | FALSE |
| CPI-SS0292 PDGFB_LRP simple:P01: simple:Q07 ENSG000000 ENSG000000 | TRUE | FALSE | TRUE |
| CPI-SS0AB2 PLXNB2_SE simple:O15 simple:Q9C ENSG000000 ENSG000000 | FALSE | TRUE | TRUE |
| CPI-SC03AF SEMA7A_in simple:O75 complex:int ENSG000000138623 | FALSE | FALSE | FALSE |
| CPI-SS0751 AGRN_NCA simple:O00: simple:P13: ENSG000000 ENSG000000 | TRUE | FALSE | TRUE |
| CPI-SS055E LAMP1_FA: simple:P11: simple:Q92 ENSG000000 ENSG000000 | TRUE | TRUE | FALSE |
| CPI-SS04F3 COPA_SOR: simple:P53: simple:Q99 ENSG000000 ENSG000000 | TRUE | FALSE | TRUE |
| CPI-CS0406 Cholesterol complex:Ch simple:P35398 ENSG000000 | TRUE | FALSE | TRUE |
| CPI-SS00F4 PTN_PTPRS simple:P21: simple:Q13 ENSG000000 ENSG000000 | TRUE | FALSE | TRUE |
| CPI-SS0715 ACKR3_CXC simple:P25: simple:P48: ENSG000000 ENSG000000 | TRUE | TRUE | FALSE |
| CPI-SS0938 DLL4_NOTC simple:Q9N simple:Q9U ENSG000000 ENSG000000 | FALSE | FALSE | TRUE |
| CPI-SS00A6 ADORA2A_ simple:P29: simple:P43: ENSG000000 ENSG000000 | TRUE | TRUE | FALSE |
| CPI-SS0C85 PROS1_AXL simple:P07: simple:P30: ENSG000000 ENSG000000 | TRUE | FALSE | TRUE |
| CPI-SS0C4D NOTCH3_JA simple:Q9U simple:Q9Y ENSG000000 ENSG000000 | FALSE | TRUE | FALSE |
| CPI-SS0B87 NOTCH1_JA simple:P46: simple:P78: ENSG000000 ENSG000000 | FALSE | TRUE | FALSE |
| CPI-SS005F UNC5B_NT: simple:Q8I: simple:Q9H ENSG000000 ENSG000000 | TRUE | TRUE | FALSE |
| CPI-SC0744 COL4A1_int simple:P02: complex:int ENSG000000187498 | TRUE | FALSE | FALSE |
| CPI-SS0593 AGRN_PTP: simple:O00: simple:Q13 ENSG000000 ENSG000000 | TRUE | FALSE | TRUE |
| CPI-SS0B99 JAG1_NOTC simple:P78: simple:Q9U ENSG000000 ENSG000000 | FALSE | FALSE | TRUE |
| CPI-SS02A0 FAM3C_CLE simple:Q92 simple:Q9U ENSG000000 ENSG000000 | TRUE | FALSE | TRUE |
| CPI-SC08F9 FN1_integri simple:P02: complex:int ENSG000000115414 | TRUE | FALSE | FALSE |
| CPI-SS02A5 LRPAP1_SO simple:P30: simple:Q99 ENSG000000 ENSG000000 | FALSE | FALSE | TRUE |
| CPI-SS0947 FLT1_VEGF: simple:P17: simple:P49: ENSG000000 ENSG000000 | TRUE | TRUE | FALSE |
| CPI-SC024F LAMC1_int: simple:P11: complex:int ENSG000000135862 | TRUE | FALSE | FALSE |
| CPI-SS07CC GRN_SORT: simple:P28: simple:Q99 ENSG000000 ENSG000000 | TRUE | FALSE | TRUE |
| CPI-SS0178 TGFB2_TGF simple:P61: simple:Q03 ENSG000000 ENSG000000 | TRUE | FALSE | TRUE |
| CPI-SC03F7 COL4A2_int simple:P08: complex:int ENSG000000134871 | TRUE | FALSE | FALSE |
| CPI-SS0DA0 TNFRSF1A_ simple:P19: simple:P28: ENSG000000 ENSG000000 | TRUE | TRUE | FALSE |

| annotation_ | Aterial EC F | Aterial EC F | Pericyte At | Pericyte At | Capillary EC | Capillary EC | Pericyte Ca | Pericyte Ca |
|-------------------------|--------------|--------------|-------------|-------------|--------------|--------------|-------------|-------------|
| curated | | 1 | | 1 | | 1 | | 1 |
| I2D,InnateDB-All | | 1 | | 1 | | 1 | | 1 |
| curated | | 1 | | 1 | | 1 | | 1 |
| curated | | 1 | | 1 | | 1 | | 1 |
| curated | | 1 | | 1 | | 1 | | 1 |
| curated | | 1 | | 1 | | 1 | | 1 |
| curated | | 1 | | 1 | | 1 | | 1 |
| curated | | 1 | | 1 | | 1 | | 1 |
| InnateDB-All | | 1 | | 1 | | 1 | | 1 |
| InnateDB-All | | 1 | | 1 | | 1 | | 1 |
| curated | | 1 | | 1 | | 1 | | 1 |
| curated | | 1 | | 1 | | 1 | | 1 |
| curated | | 1 | | 1 | | 1 | | 1 |
| curated | | 1 | | 1 | | 1 | | 1 |
| curated | | 1 | | 1 | | 1 | | 1 |
| curated | | 1 | | 1 | | 1 | | 1 |
| curated | | 1 | 0.199 | 0 | | 1 | | 1 |
| InnateDB-All | | 1 | | 0.963 | | 1 | | 0.884 |
| curated | | 1 | | 1 | | 1 | | 1 |
| curated | | 1 | | 1 | | 1 | | 1 |
| curated | | 1 | | 1 | | 1 | | 1 |
| curated | | 1 | | 1 | | 1 | | 1 |
| curated | | 1 | | 1 | | 1 | | 1 |
| curated | | 1 | | 1 | | 1 | | 1 |
| curated | | 1 | | 1 | 0.603 | | | 1 |
| curated | | 1 | | 1 | | 1 | | 1 |
| curated | | 1 | | 0.985 | | 1 | | 1 |
| curated | | 0.204 | | 1 | | 1 | | 1 |
| curated | | 0.899 | | 1 | | 1 | | 1 |
| curated | | 1 | | 1 | | 1 | | 1 |
| curated | | 1 | | 1 | | 1 | | 1 |
| curated | | 1 | | 0.393 | | 1 | | 1 |
| I2D,IMEx,InnateDB-All,I | | 1 | 3.959 | 0 | | 1 | | 1 |
| curated | | 1 | | 1 | | 1 | | 1 |
| curated | 14.431 | 0.038 | | 1 | | 1 | | 1 |
| curated | | 1 | 0.179 | 0 | | 1 | | 1 |
| curated | | 1 | | 1 | | 1 | | 1 |
| curated | | 1 | | 1 | | 1 | | 1 |
| curated | 0.225 | 0.008 | | 1 | | 0.29 | | 1 |
| curated | | 0.928 | | 1 | | 1 | | 1 |
| curated | | 1 | | 0.961 | | 1 | | 0.546 |
| curated | | 0.244 | | 1 | | 0.962 | | 1 |
| curated | | 1 | 1.872 | 0 | | 1 | 1.805 | 0 |
| curated | | 1 | 7.901 | 0 | | 1 | | 1 |
| curated | | 1 | | 1 | | 1 | | 1 |

| | | | | | | | | |
|------------------------------------|-------|-------|-------|-------|-------|-------|-------|-------|
| curated | | 0.576 | | 1 | | 1 | | 1 |
| curated | | 1 | 0.845 | 0 | | 1 | | 0.338 |
| curated | 0.625 | 0 | | 1 | | 1 | | 1 |
| curated | 3.508 | 0.04 | | 1 | | 1 | | 1 |
| curated | | 1 | 1.315 | 0 | | 1 | | 1 |
| curated | 3.584 | 0.024 | | 1 | | 1 | | 1 |
| curated | | 1 | | 0.211 | | 1 | | 1 |
| curated | 2.571 | 0 | | 1 | 2.45 | 0 | | 1 |
| curated | | 1 | 0.787 | 0 | | 1 | | 1 |
| curated | | 1 | | 1 | | 1 | | 1 |
| I2D,InnateC | 1.298 | 0 | | 1 | 1.177 | 0.04 | | 1 |
| curated | | 1 | | 0.467 | | 1 | | 0.837 |
| curated | | 1 | | 1 | | 1 | | 1 |
| curated | | 0.657 | | 1 | | 1 | | 1 |
| InnateDB-All | | 1 | | 1 | | 1 | | 1 |
| IMEx,InnateDB-All,IntAc | | 0.71 | | 1 | | 1 | | 1 |
| curated | | 1 | | 1 | | 1 | | 1 |
| I2D,IMEx,In | 3.944 | 0 | | 1 | | 1 | | 1 |
| guidetopharmacology.c | | 1 | | 1 | | 1 | | 0.992 |
| curated | 1.442 | 0 | | 1 | 1.226 | 0 | | 1 |
| I2D,IMEx,InnateDB-All,IntAc | | 1 | 0.405 | 0 | | 1 | 0.345 | 0.039 |
| guidetopha | 0.487 | 0 | | 1 | | 0.221 | | 1 |
| curated | | 1 | 1.672 | 0 | | 1 | 1.363 | 0 |
| curated | | 0.961 | | 1 | | 1 | | 1 |
| curated | 0.58 | 0 | | 1 | | 0.776 | | 1 |
| curated | | 1 | | 1 | | 1 | | 1 |
| curated | | 0.15 | | 1 | | 1 | | 1 |
| curated | 1.419 | 0 | | 1 | 1.231 | 0 | | 1 |
| IMEx,InnateDB-All,IntAc | | 1 | | 0.796 | | 1 | | 1 |
| curated | 3.508 | 0.04 | | 1 | | 1 | | 1 |
| curated | | 0.906 | | 1 | | 1 | | 1 |
| curated | | 0.054 | | 1 | | 1 | | 1 |
| curated | | 1 | | 1 | | 1 | | 1 |
| IMEx,InnateDB-All,MIN ⁺ | | 0.814 | | 1 | | 1 | | 1 |
| curated | 0.841 | 0 | | 1 | | 1 | | 1 |
| curated | | 1 | | 1 | | 1 | | 1 |
| IMEx,InnateDB-All,MIN ⁺ | | 1 | | 1 | | 1 | | 1 |

| Venous EC | Venous EC | Pericyte | Ve Pericyte | Ve EC-C4 | Peric EC-C4 | Peric Pericyte | EC Pericyte | EC-C4_qual |
|-----------|-----------|----------|-------------|----------|-------------|----------------|-------------|------------|
| | 1 | | 1 | | 1 | 0.898 | 0 | |
| 0.556 | 0 | | 1 | | 1 | | 1 | |
| | 1 | | 1 | 0.298 | 0 | | 1 | |
| 1.382 | 0 | | 1 | | 0.992 | | 1 | |
| 2.183 | 0 | | 1 | | 0.111 | | 1 | |
| | 1 | 0.226 | 0 | | 1 | | 1 | |
| 0.178 | 0 | | 1 | | 1 | | 1 | |
| 0.364 | 0 | | 1 | | 1 | | 1 | |
| | 1 | | 1 | 0.438 | 0 | | 1 | |
| | 1 | | 1 | 1.686 | 0 | | 1 | |
| | 1 | 2.808 | 0 | | 1 | 2.034 | 0 | |
| | 1 | 2.817 | 0 | | 1 | 2.043 | 0 | |
| | 1 | 2.808 | 0 | | 1 | 2.034 | 0 | |
| | 1 | 2.815 | 0 | | 1 | 2.041 | 0 | |
| | 1 | 2.829 | 0 | | 1 | 2.055 | 0 | |
| | 1 | 2.813 | 0 | | 1 | 2.039 | 0 | |
| | 1 | 0.21 | 0 | | 1 | | 1 | |
| | 1 | 0.473 | 0 | | 1 | 0.404 | 0 | |
| 0.185 | 0 | | 1 | 0.158 | 0 | | 1 | |
| 0.214 | 0 | | 1 | 0.185 | 0 | | 1 | |
| 0.185 | 0 | | 1 | 0.158 | 0 | | 1 | |
| 0.214 | 0 | | 1 | 0.185 | 0 | | 1 | |
| 0.532 | 0 | | 1 | 0.429 | 0 | | 1 | |
| 1.375 | 0 | | 1 | | 1 | | 1 | |
| 0.247 | 0 | 0.401 | 0 | | 1 | 0.285 | 0 | |
| | 0.265 | | 1 | 2.327 | 0 | | 1 | |
| 50.255 | 0 | | 1 | | 1 | | 1 | |
| | 1 | 2.851 | 0 | | 1 | 2.077 | 0 | |
| 0.412 | 0 | 0.309 | 0 | 0.227 | 0 | | 1 | |
| | 1 | 6.75 | 0 | | 1 | 6.322 | 0 | |
| | 1 | 4.66 | 0 | | 1 | | 1 | |
| 0.202 | 0 | 0.277 | 0 | | 1 | 0.238 | 0 | |
| 21.571 | 0 | | 1 | 14.707 | 0 | | 1 | |
| | 1 | 0.171 | 0 | | 1 | 0.217 | 0 | |
| 2.176 | 0 | | 1 | | 0.543 | | 1 | |
| 0.664 | 0 | 0.664 | 0 | | 0.556 | | 0.556 | |
| 0.319 | 0 | | 1 | 0.248 | 0 | | 1 | |
| 1.438 | 0 | 0.71 | 0.023 | 0.864 | 0 | | 1 | |
| | 0.062 | 0.309 | 0 | | 0.664 | 0.254 | 0.039 | |
| 1.123 | 0 | | 1 | 1.259 | 0 | | 1 | |
| | 1 | 1.879 | 0 | | 1 | 1.887 | 0 | |
| 14.076 | 0 | 8.106 | 0 | | 1 | 7.979 | 0 | |
| 0.365 | 0 | 0.365 | 0 | 0.217 | 0 | 0.217 | 0 | |

| | | | | | | | |
|--------|-------|--------|-------|-------|-------|--------|-------|
| 1.075 | 0 | | 1 | 1.211 | 0 | | 1 |
| | 0.605 | 1.088 | 0 | | 1 | | 0.22 |
| 0.754 | 0 | 0.959 | 0 | 0.659 | 0 | | 0.976 |
| 4.894 | 0 | | 1 | 4.755 | 0 | | 1 |
| | 1 | 1.738 | 0 | | 1 | 1.199 | 0 |
| 4.97 | 0 | | 1 | 4.831 | 0 | | 1 |
| | 1 | 0.88 | 0 | | 1 | | 1 |
| 3.082 | 0 | | 1 | 2.666 | 0 | | 1 |
| | 1 | 0.643 | 0 | | 1 | 0.726 | 0 |
| | 1 | | 0.244 | | 1 | 1.141 | 0 |
| 1.808 | 0 | | 1 | 1.393 | 0 | | 1 |
| | 1 | 0.3 | 0 | | 1 | 0.254 | 0 |
| | 0.97 | | 1 | 2.382 | 0 | | 1 |
| 0.717 | 0 | | 1 | | 0.068 | | 1 |
| 1.98 | 0 | | 1 | 1.328 | 0.031 | | 1 |
| 0.745 | 0 | | 0.392 | 0.549 | 0.015 | | 1 |
| | 0.945 | | 1 | 1.146 | 0 | | 1 |
| 4.645 | 0 | | 1 | | 1 | | 1 |
| | 1 | 15.364 | 0 | | 1 | 12.077 | 0.008 |
| 1.433 | 0 | | 1 | 1.255 | 0 | | 1 |
| | 1 | 0.548 | 0 | | 1 | 0.437 | 0 |
| 0.666 | 0 | 0.604 | 0 | 0.648 | 0 | | 1 |
| | 1 | 1.694 | 0 | | 1 | 1.593 | 0 |
| 2.336 | 0 | | 1 | 1.501 | 0 | | 1 |
| 0.694 | 0 | 0.427 | 0 | | 0.209 | | 1 |
| 3.662 | 0 | 3.555 | 0 | | 0.101 | | 0.467 |
| 0.743 | 0 | | 1 | 0.568 | 0 | | 1 |
| 1.654 | 0 | 2.124 | 0 | 1.414 | 0 | | 0.704 |
| | 1 | 2.351 | 0 | | 1 | 2.594 | 0 |
| 4.894 | 0 | | 1 | 4.755 | 0 | | 1 |
| 0.614 | 0 | 0.473 | 0.016 | 0.483 | 0.008 | | 1 |
| 21.564 | 0 | | 1 | 14.7 | 0 | | 1 |
| | 1 | 2.334 | 0 | | 1 | 2.052 | 0 |
| 0.637 | 0 | | 0.091 | 0.578 | 0 | | 1 |
| 0.943 | 0 | | 1 | | 0.471 | | 1 |
| 2.627 | 0 | 3.121 | 0 | 2.524 | 0.008 | | 0.164 |
| | 0.392 | 0.595 | 0 | 0.723 | 0 | 0.536 | 0.024 |