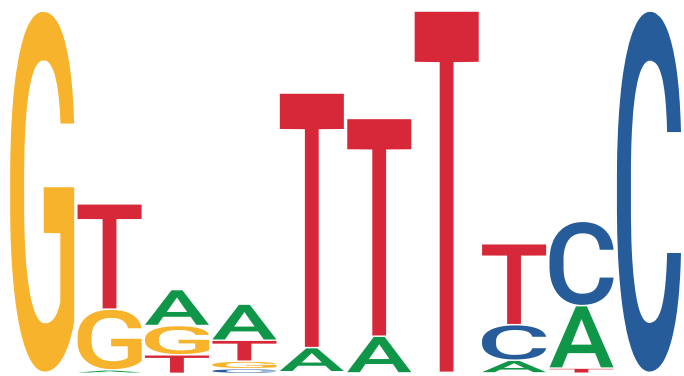


nvec_dap_1
Nvec_vc1.1_XM_048733736.1
OG: RHD_Rel.HG1.0:like:NFAT5/NFATC1/NFATC2/NFATC3/NFATC4
PFAM: RHD_DNA_bind/RHD_dimer

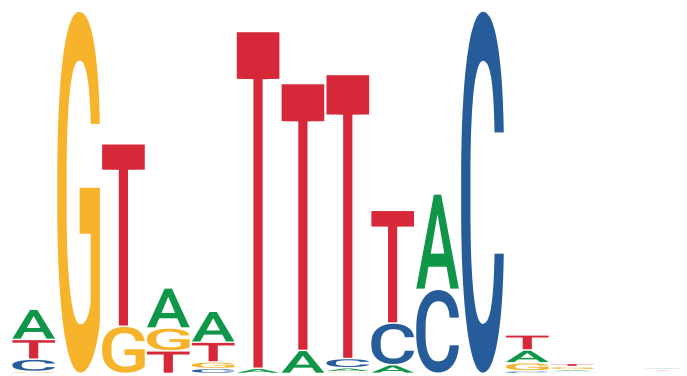
dimont



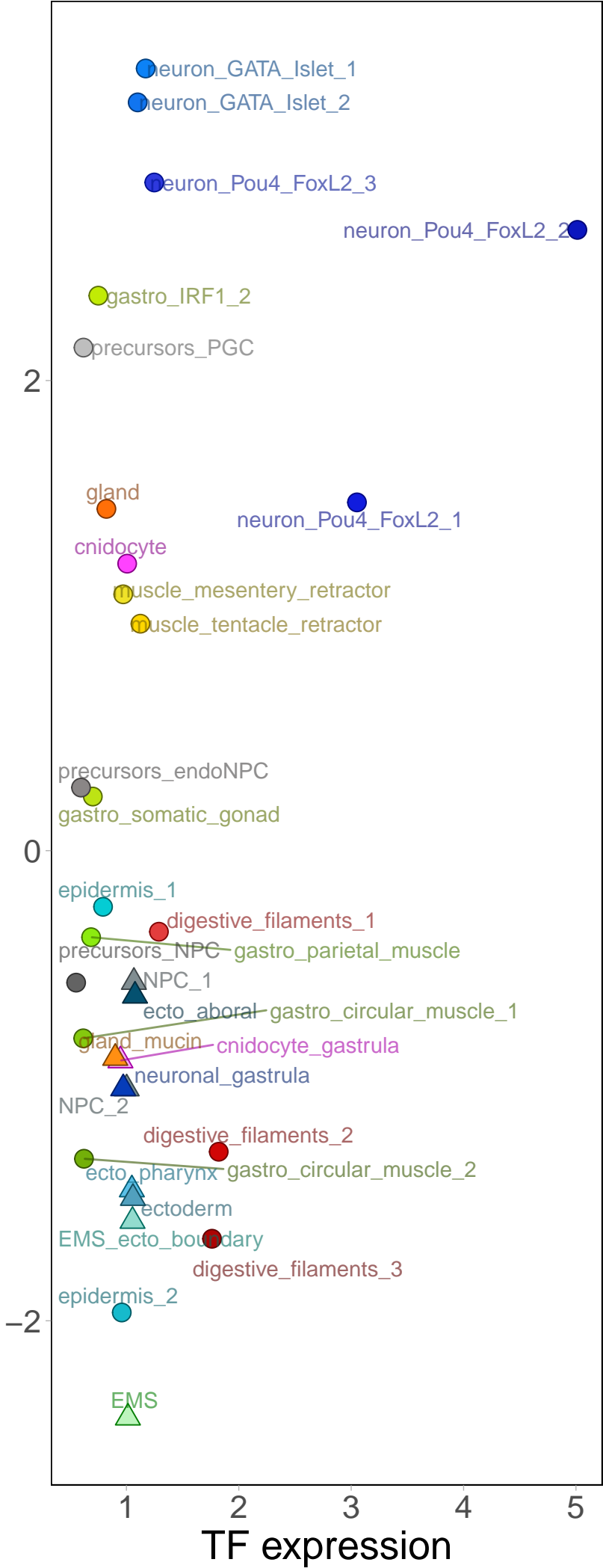
homer



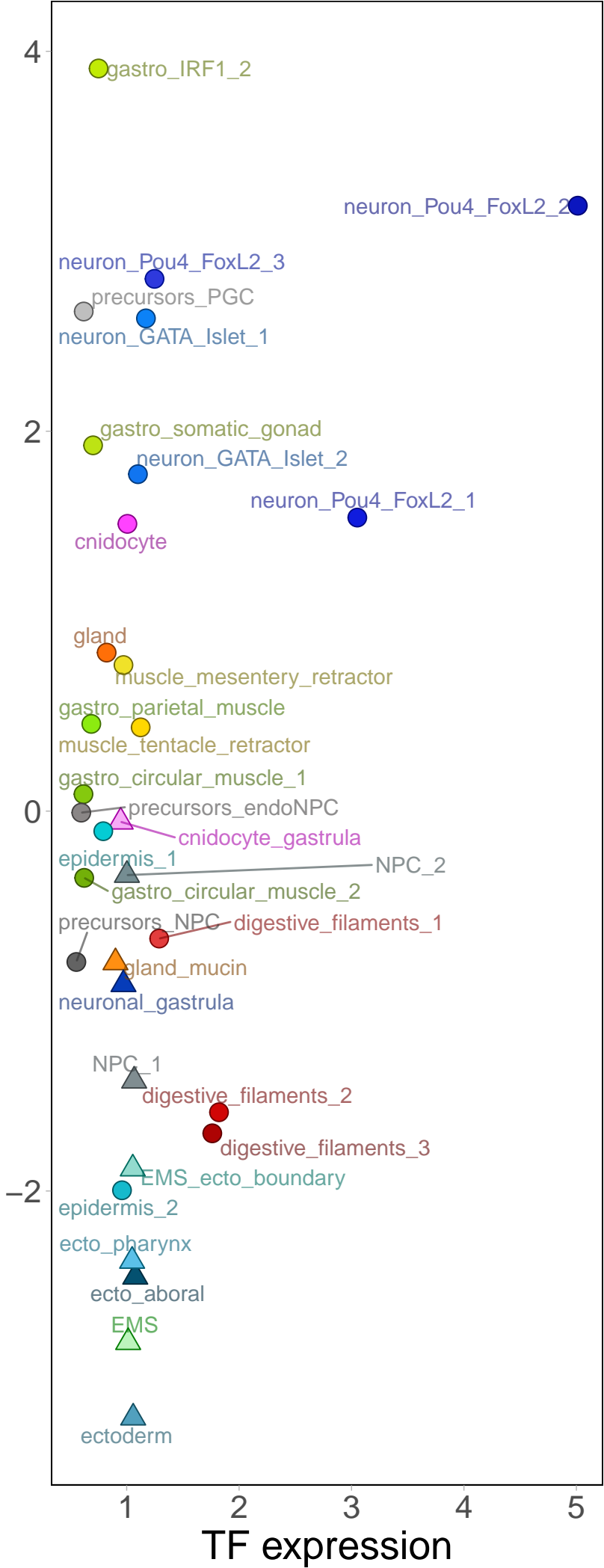
streme



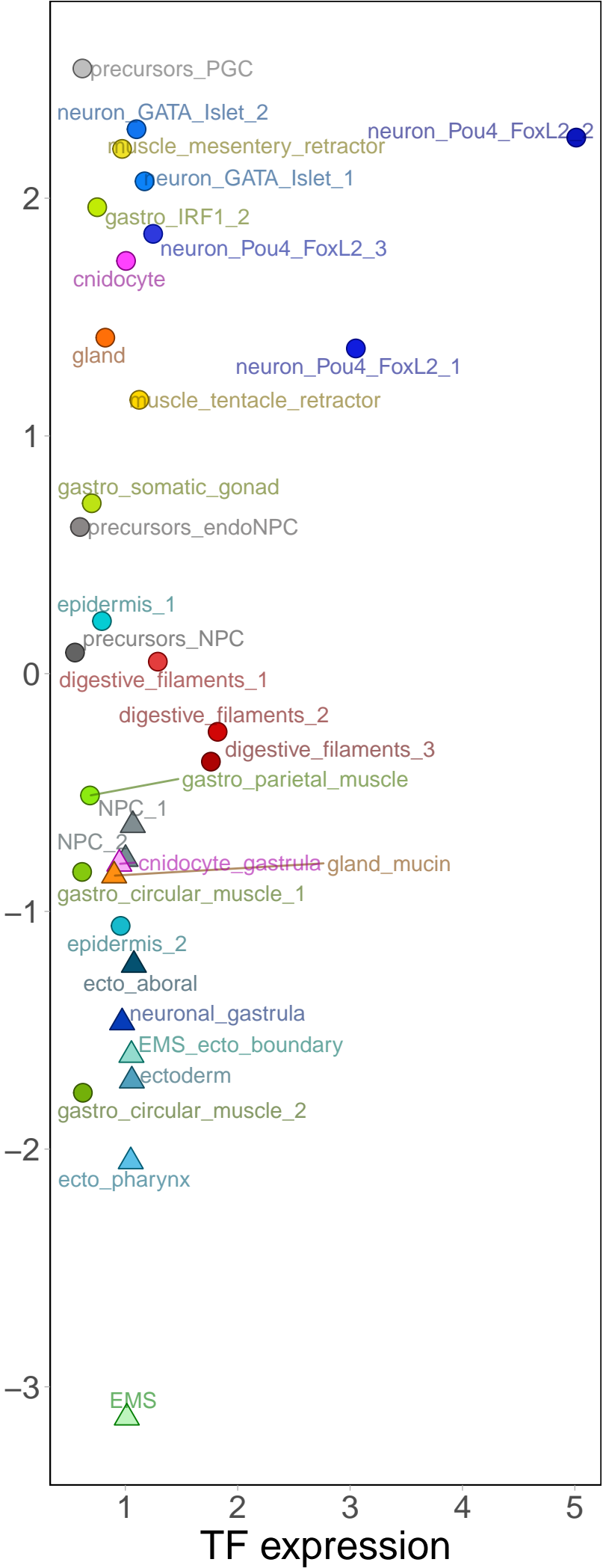
TF activity



TF activity



TF activity



nvec_dap_2
Nvec_vc1.1_XM_032382466.2
OG: bZIP.HG2.5:CEBPA/CEBPB/CEBPD/CEBPE/CEBPG
PFAM: bZIP_2

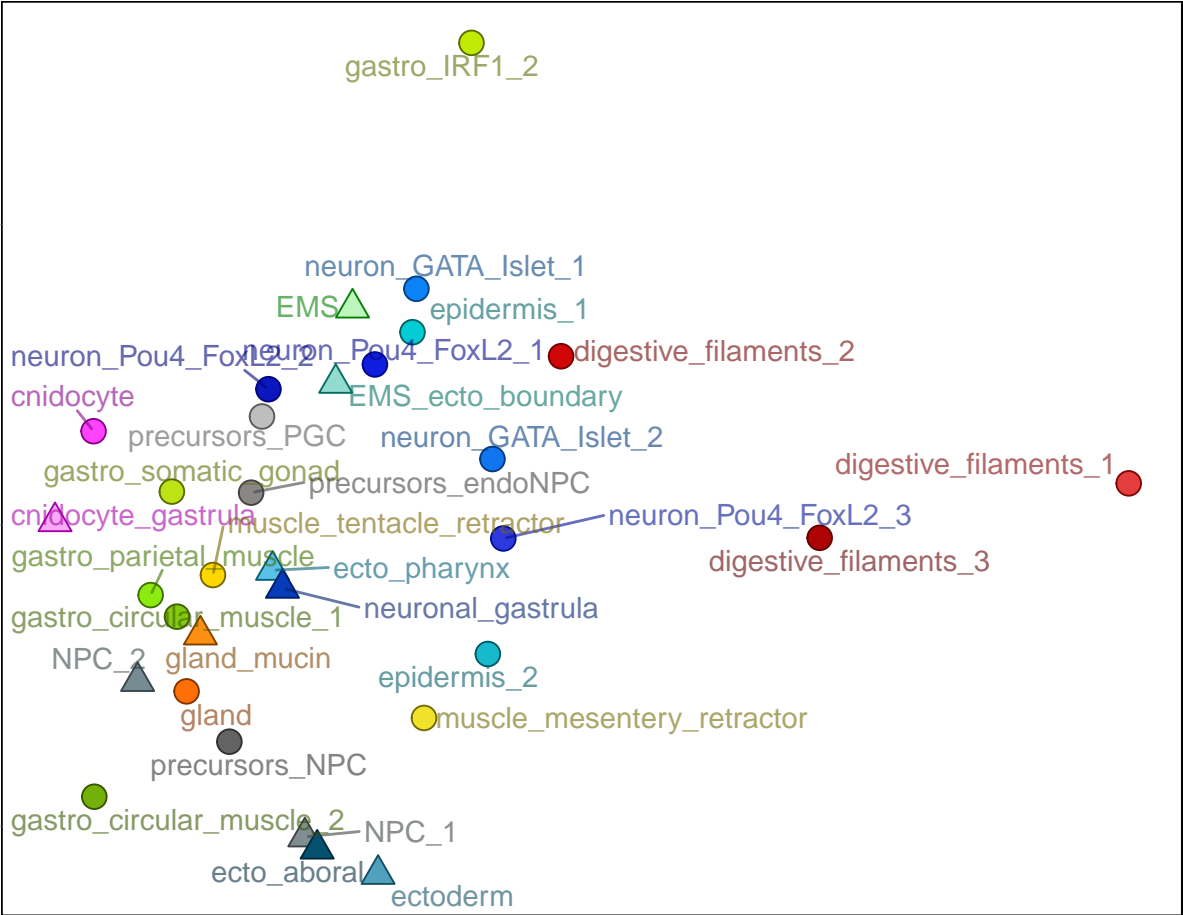
archetype



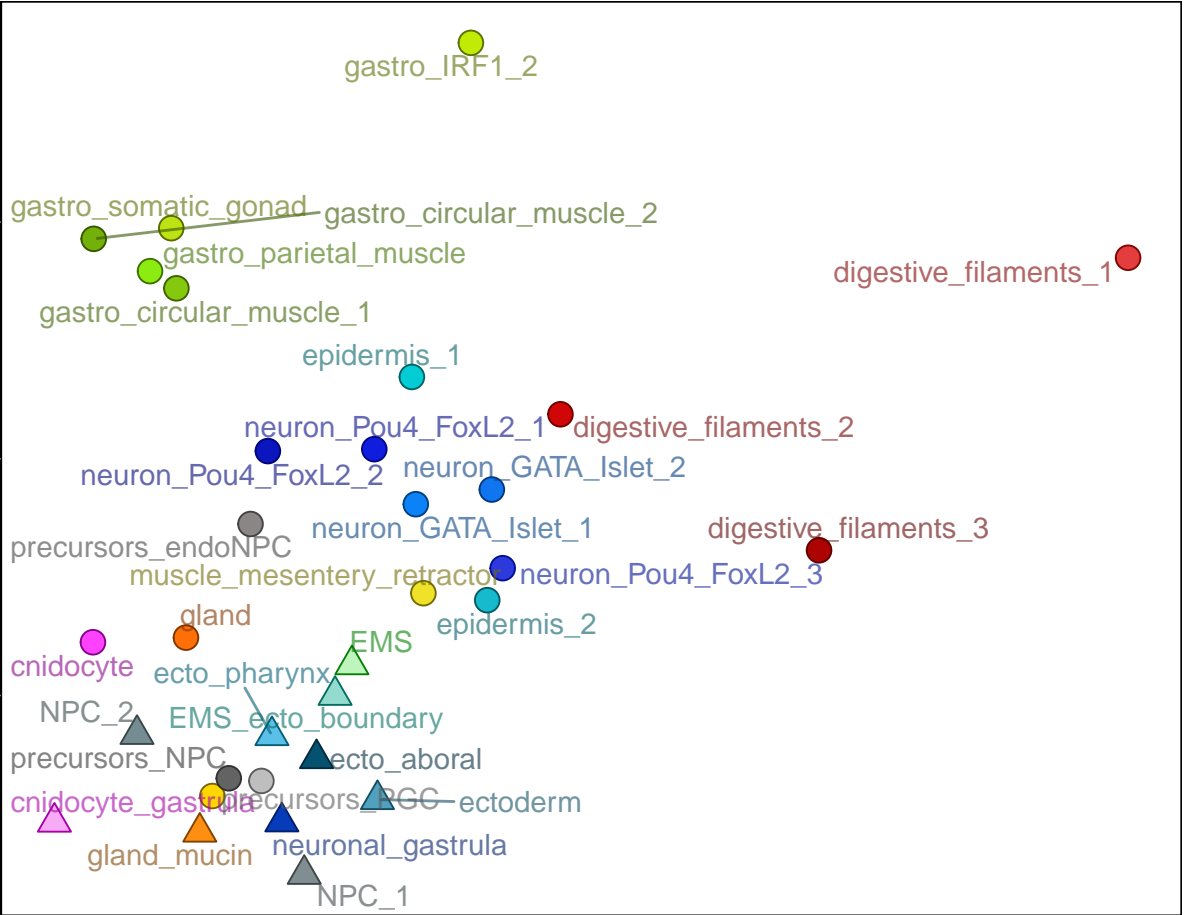
dimont



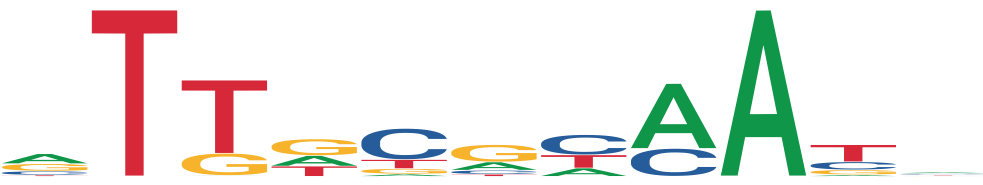
TF activity



TF activity



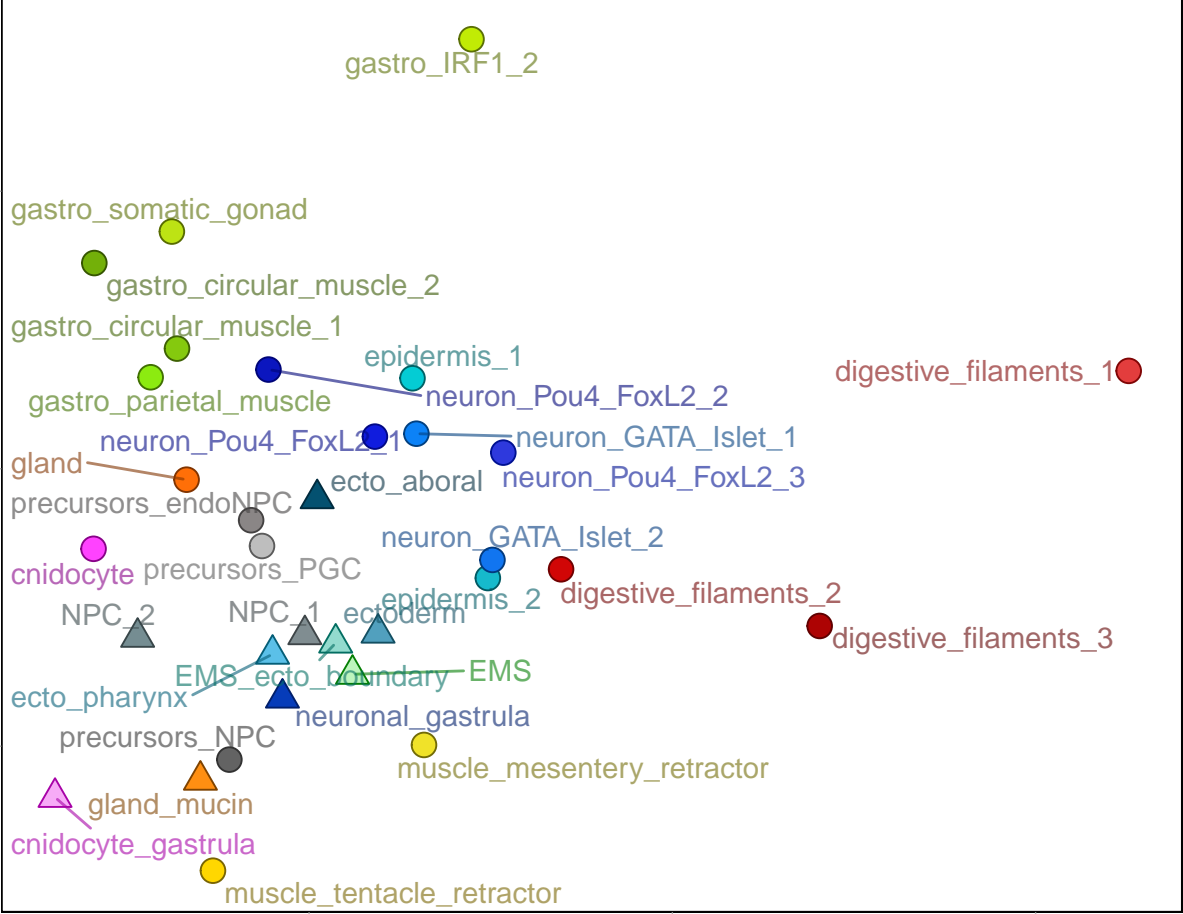
homer



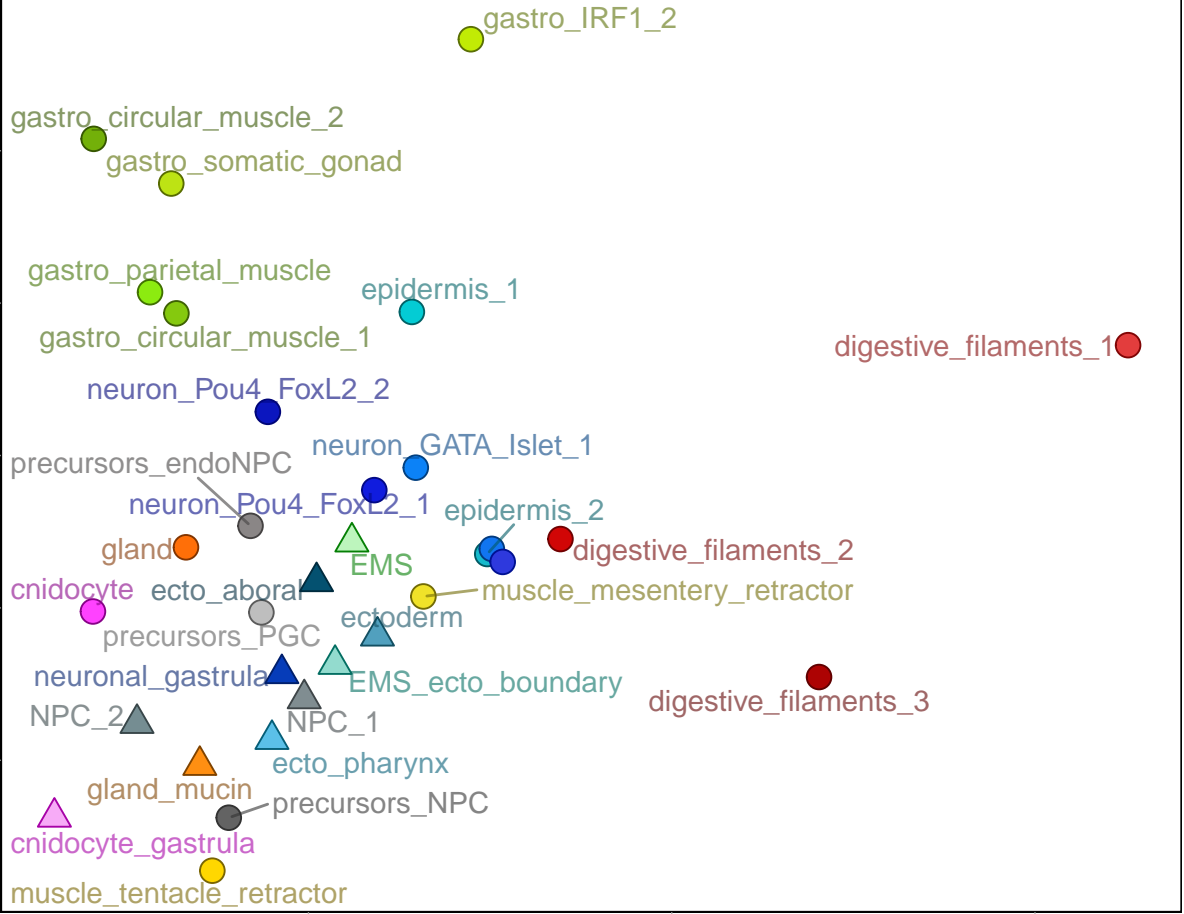
streame



TF activity



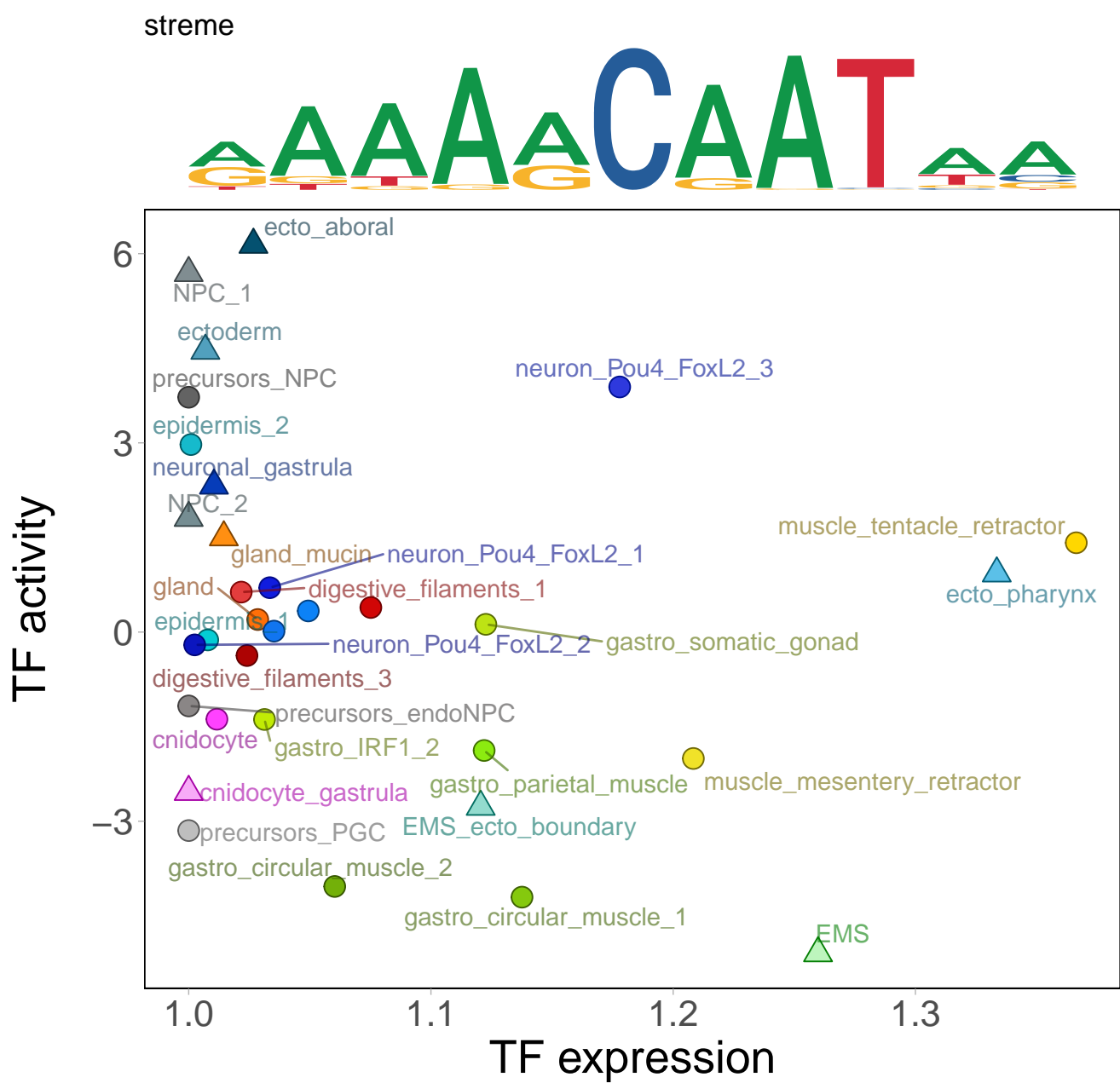
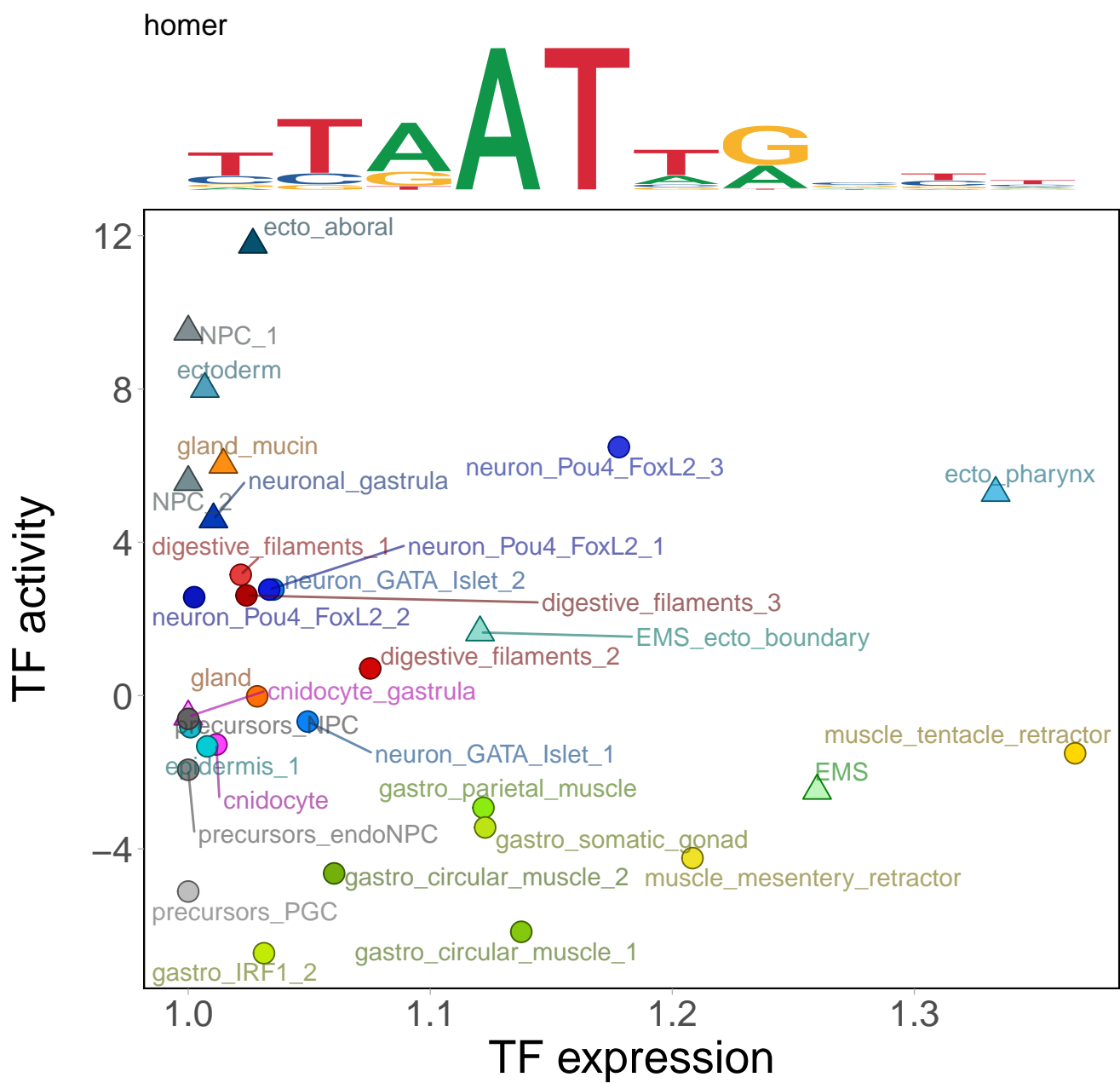
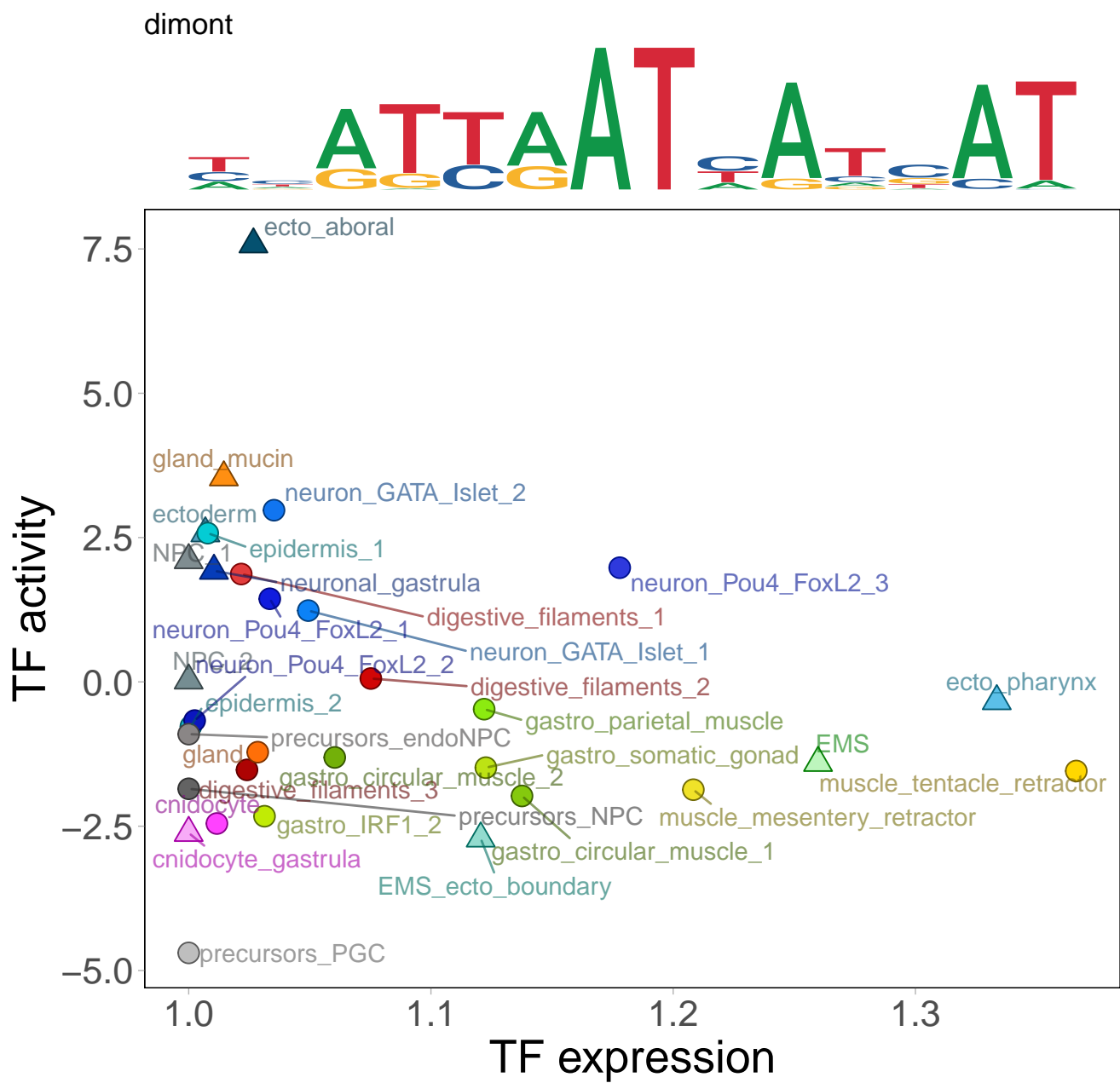
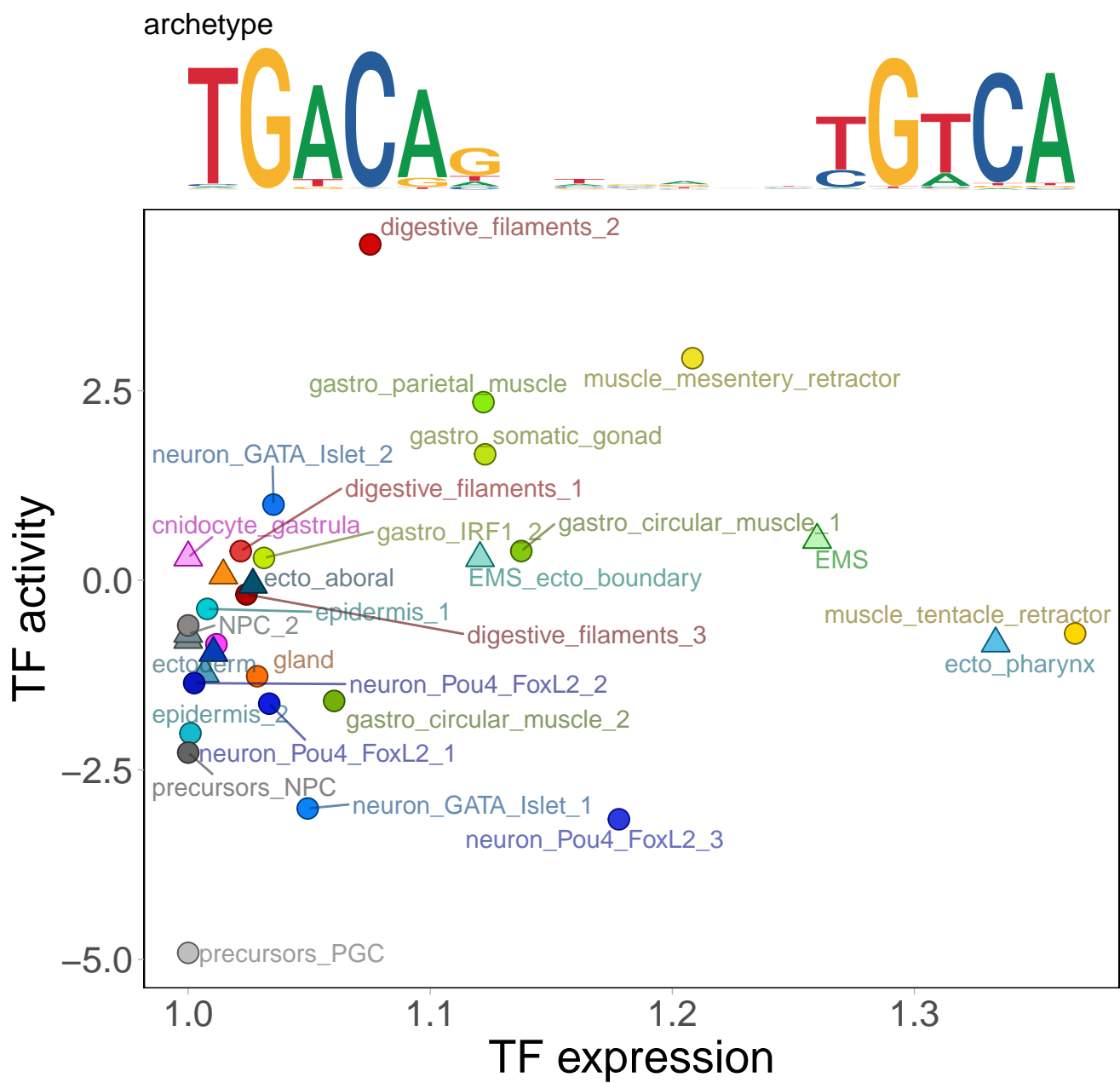
TF activity



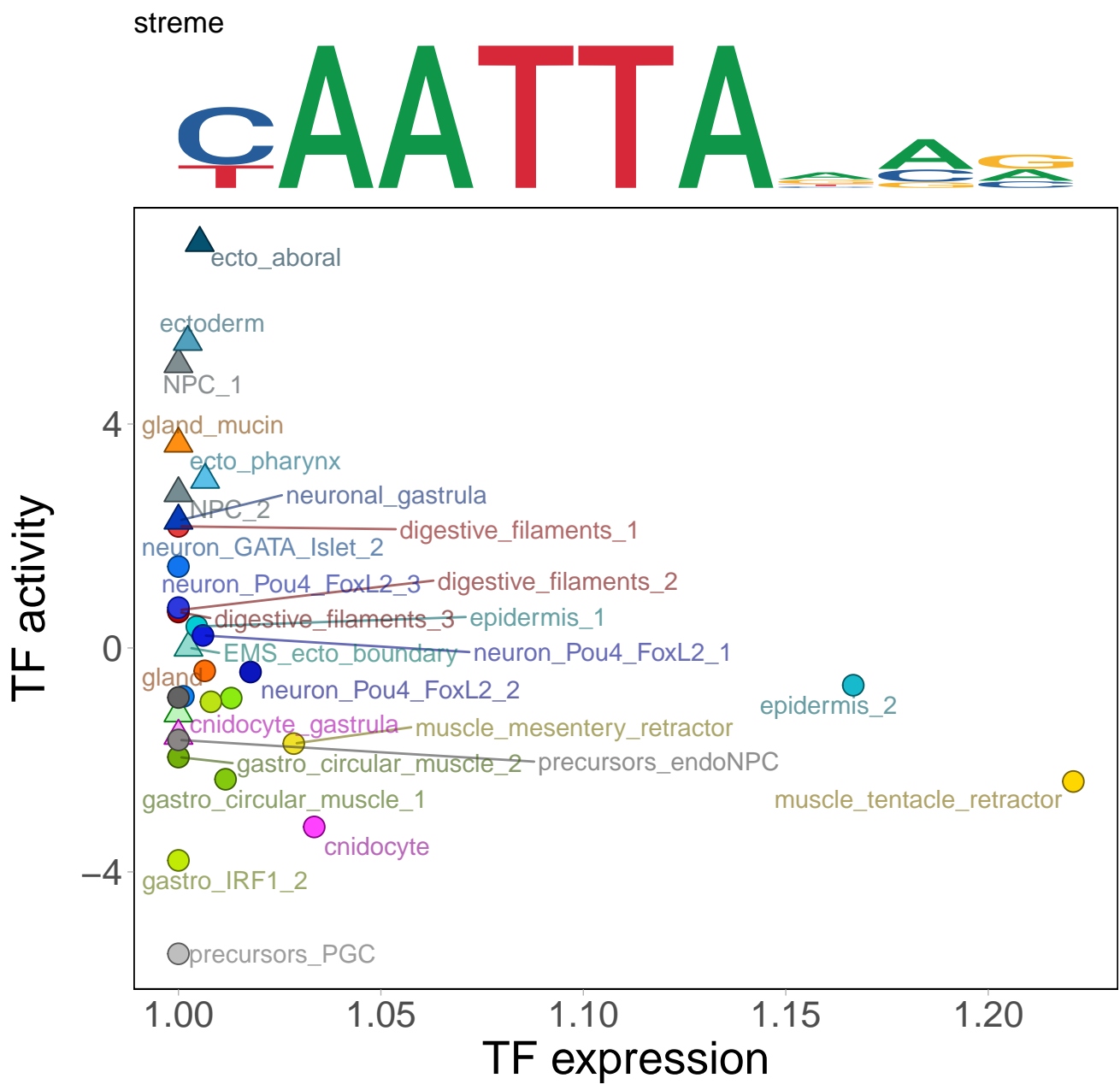
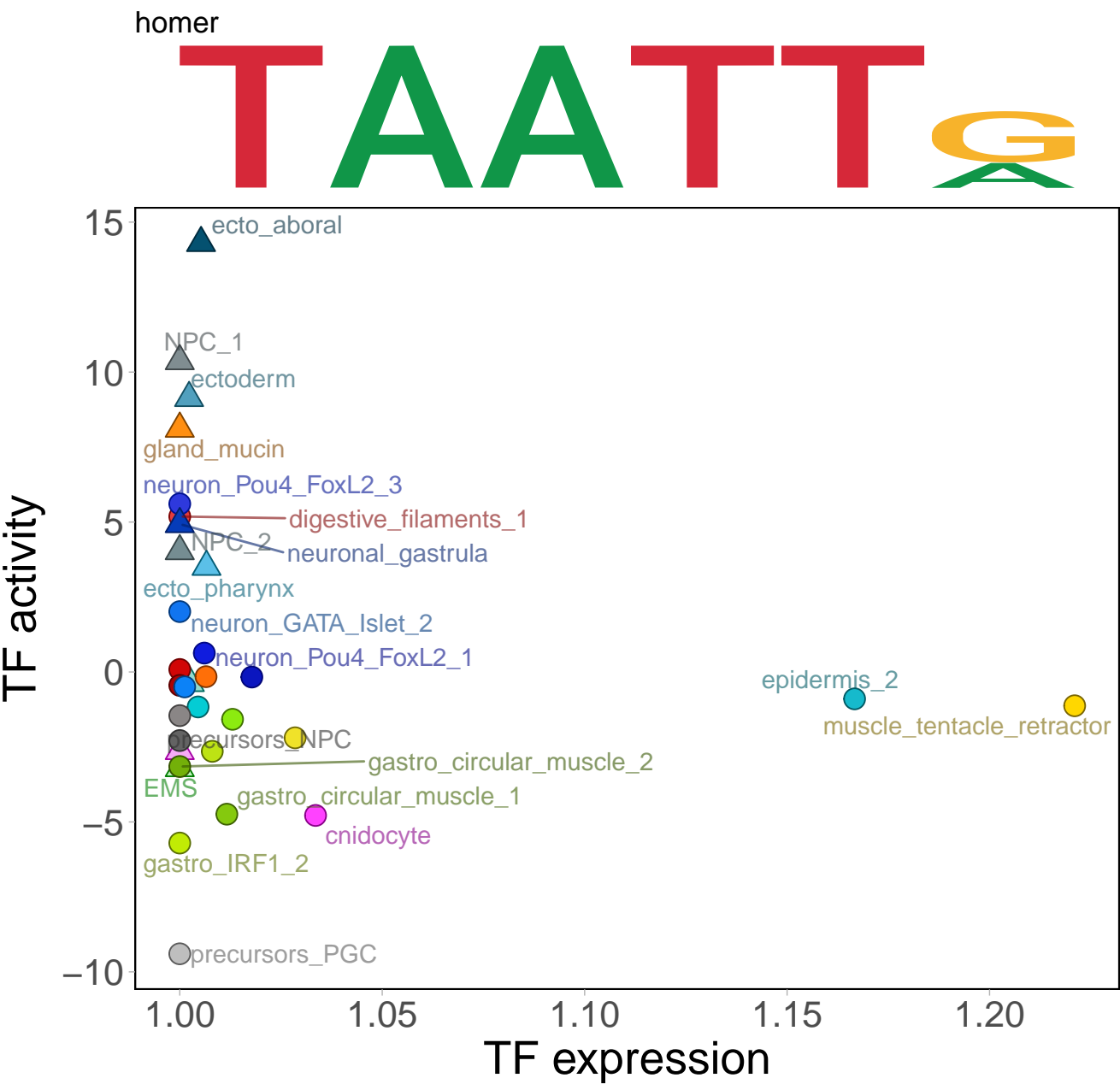
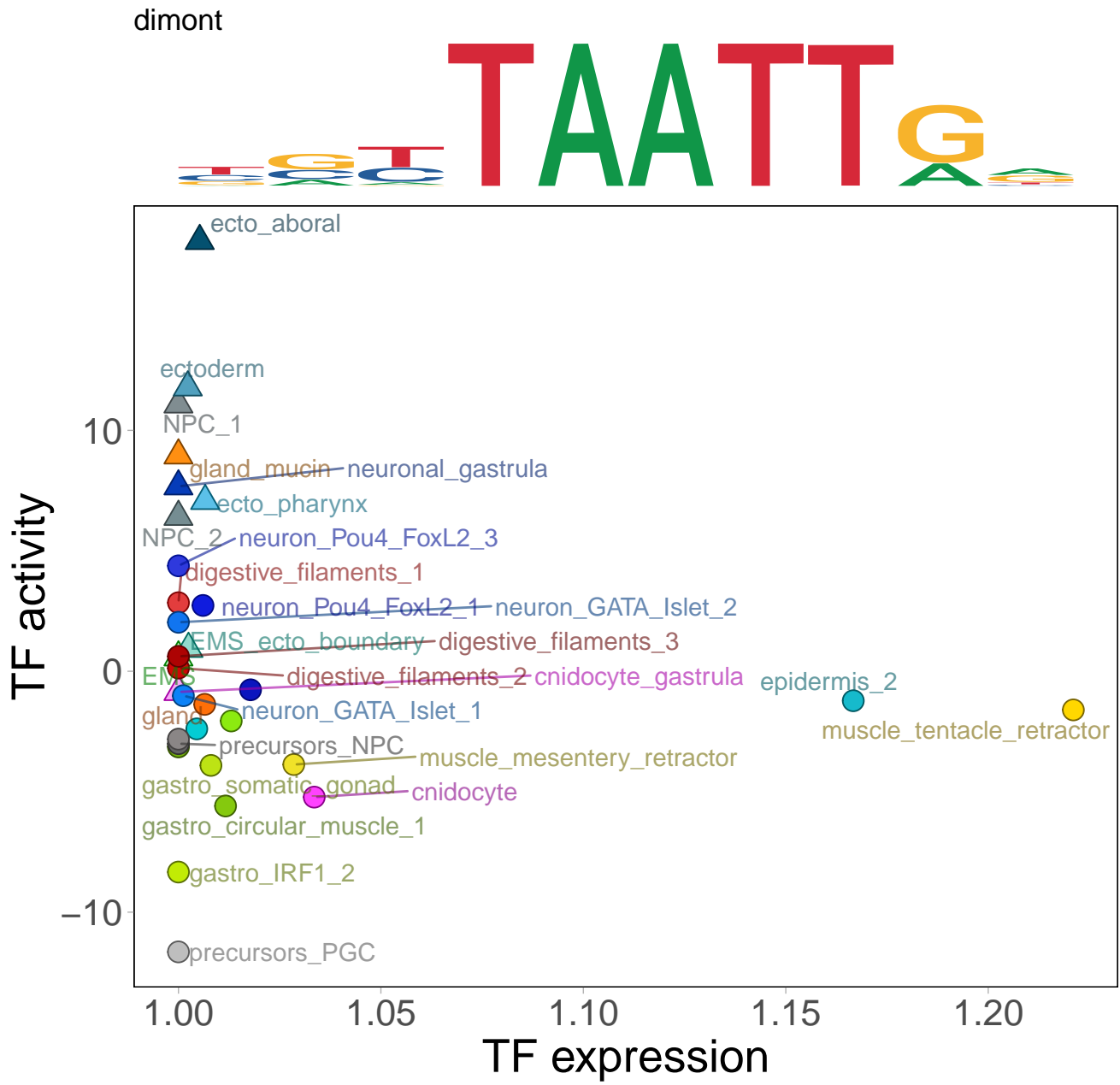
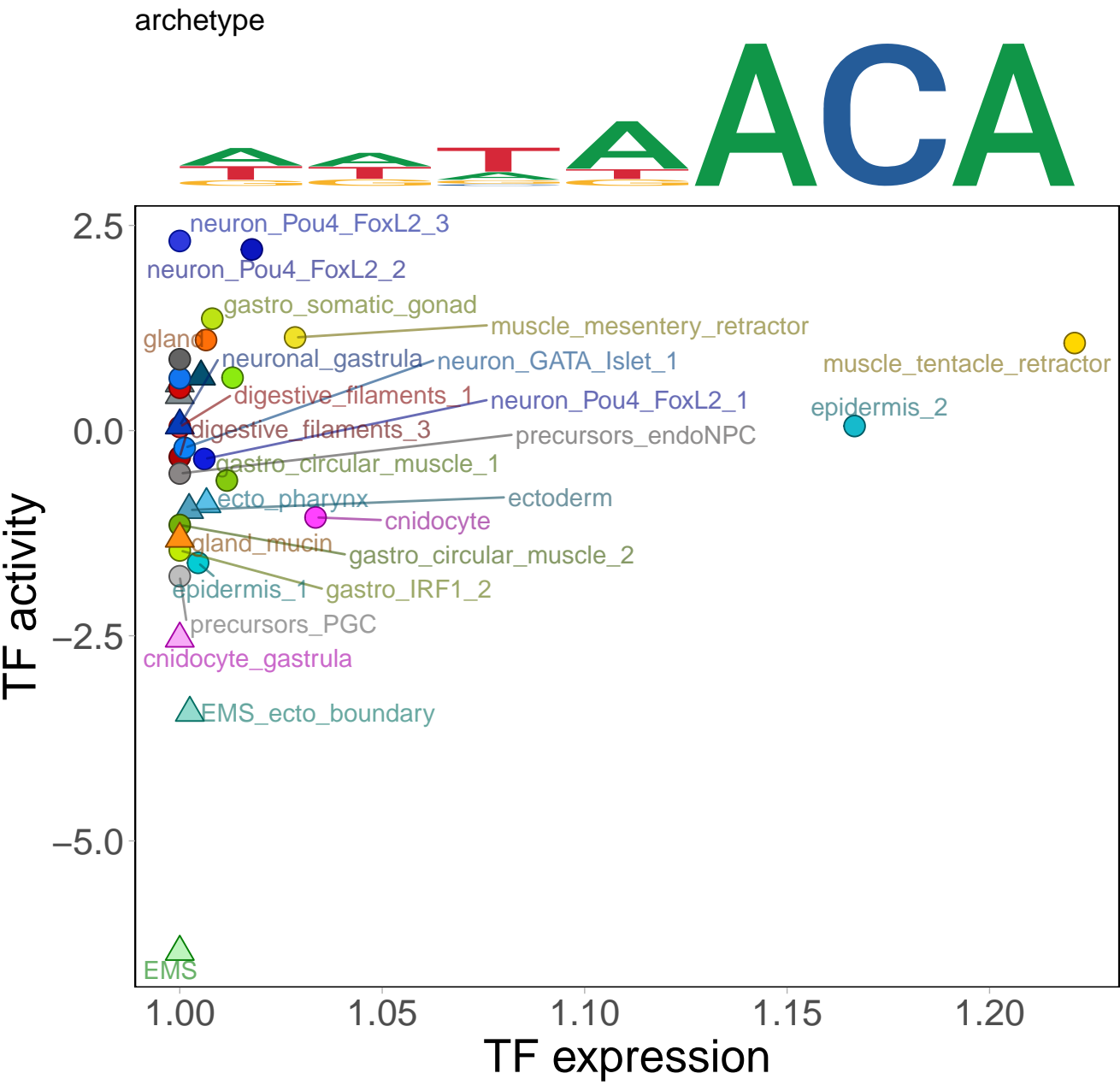
TF expression

TF expression

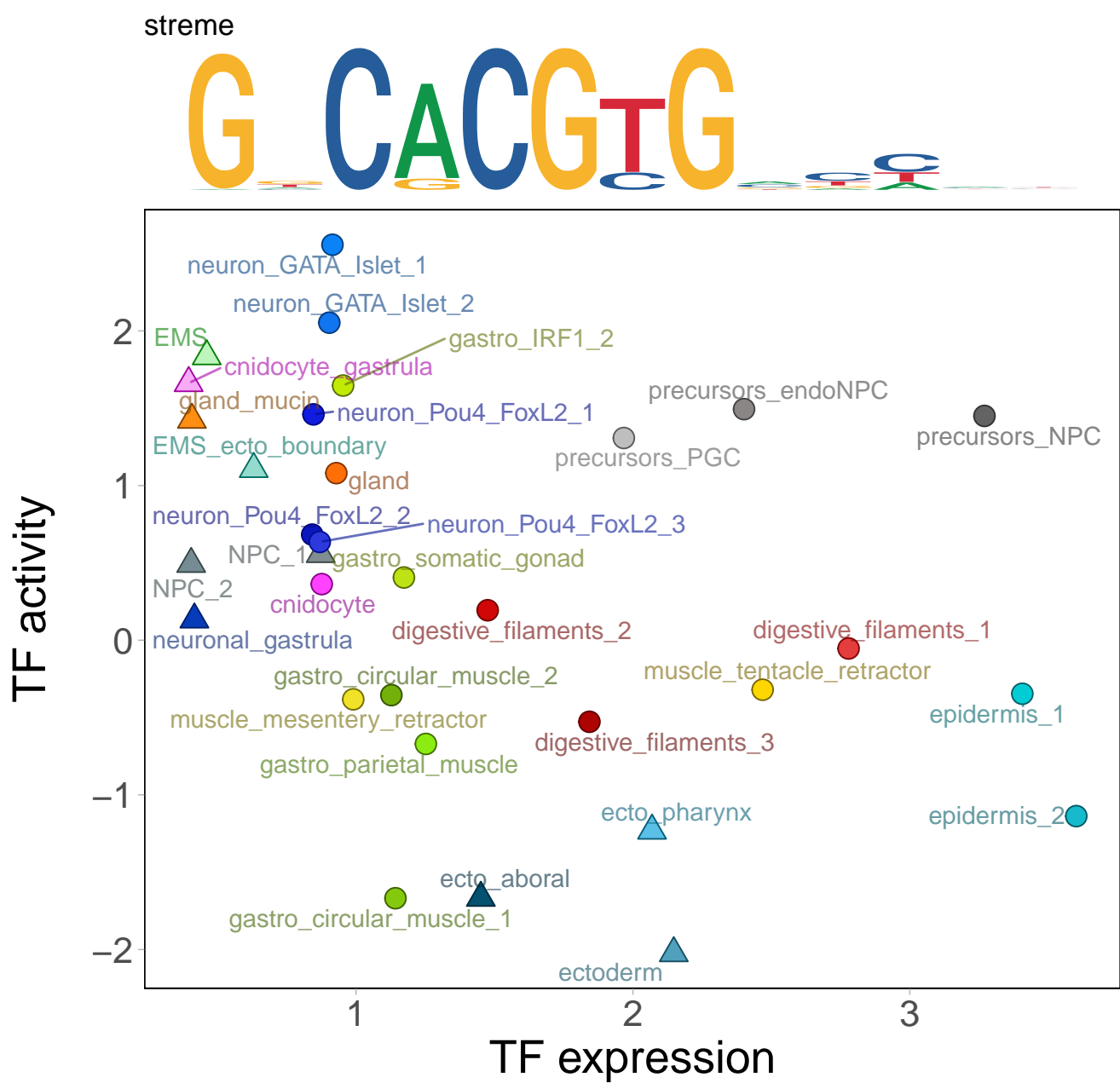
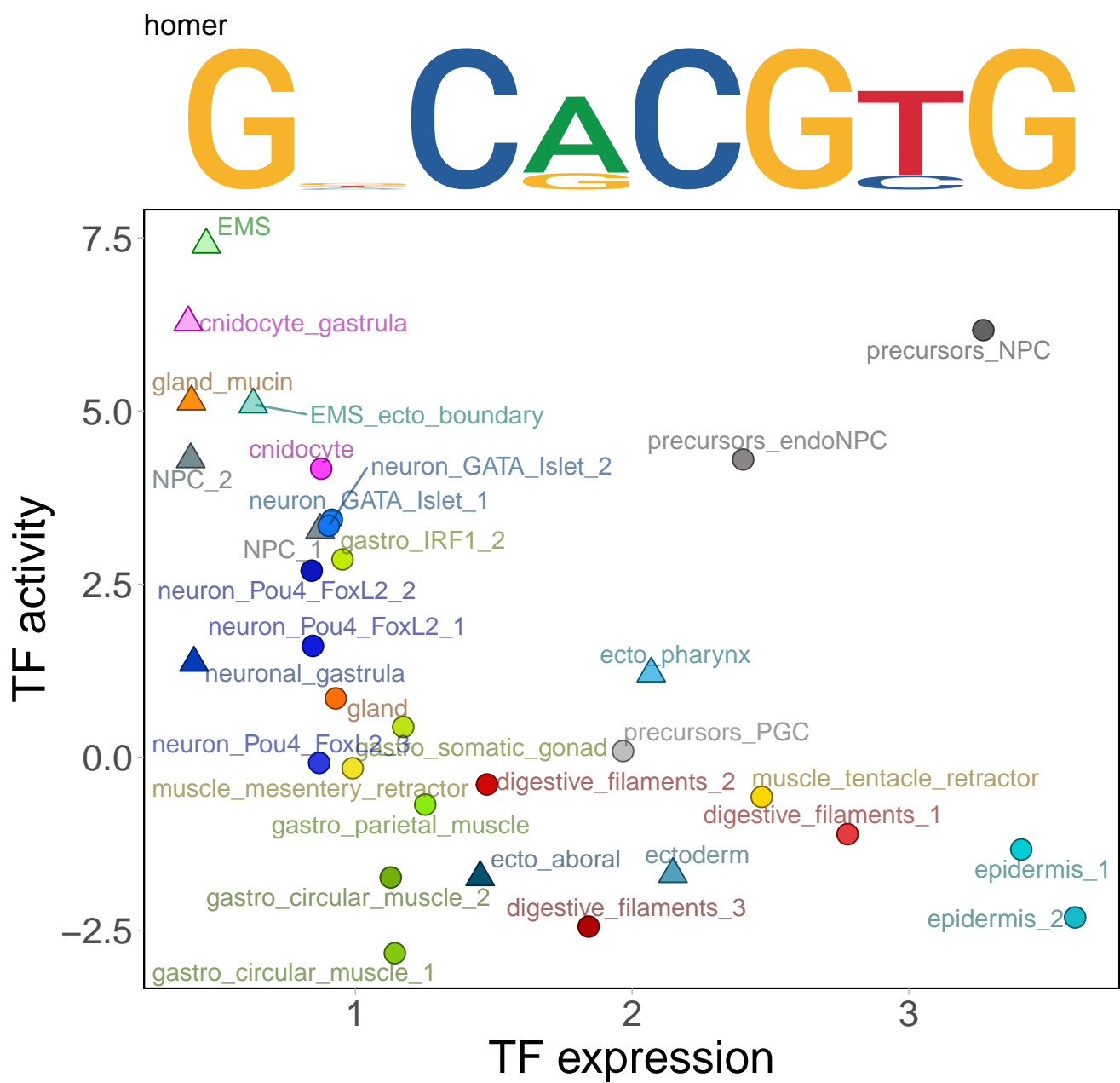
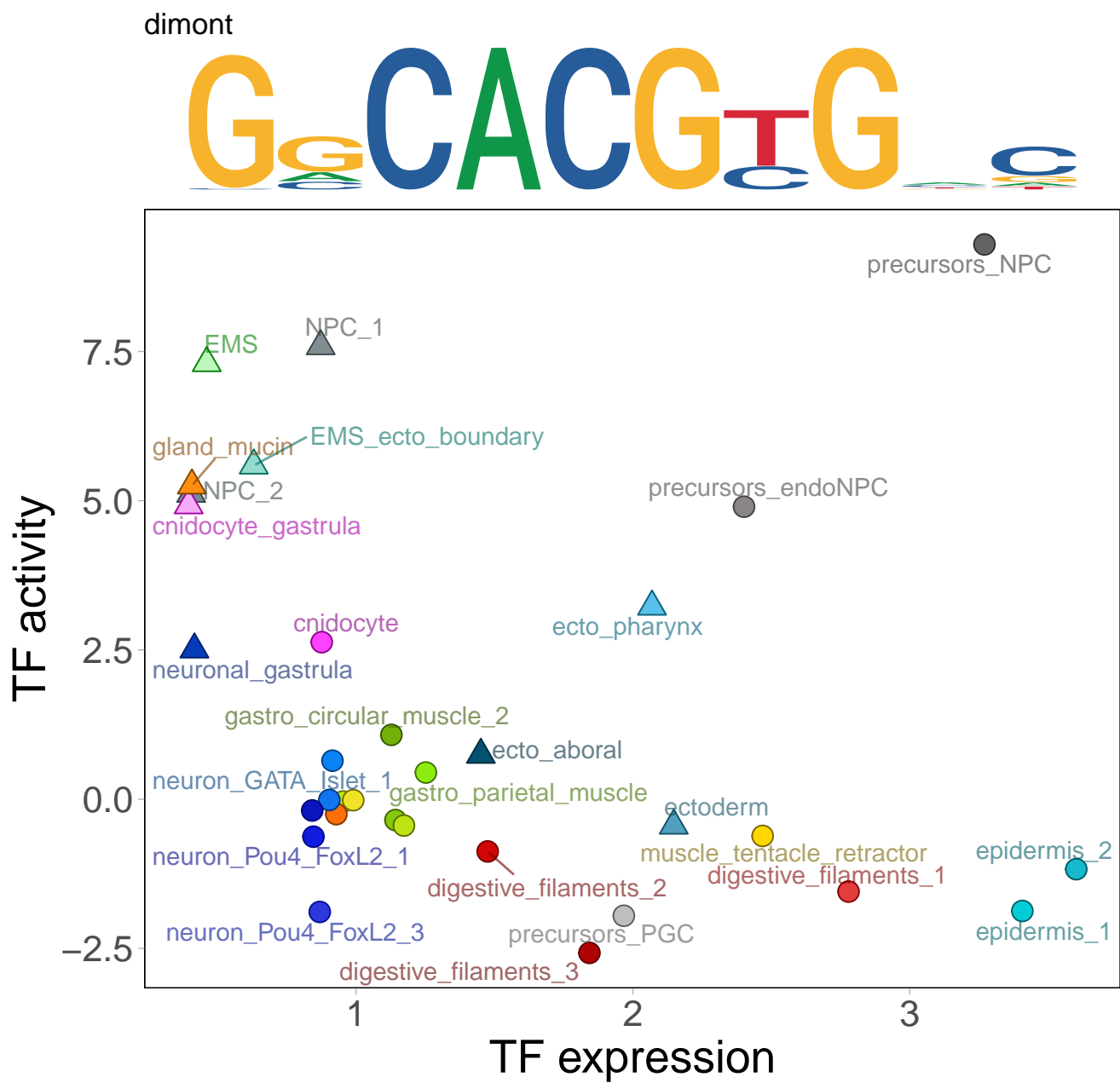
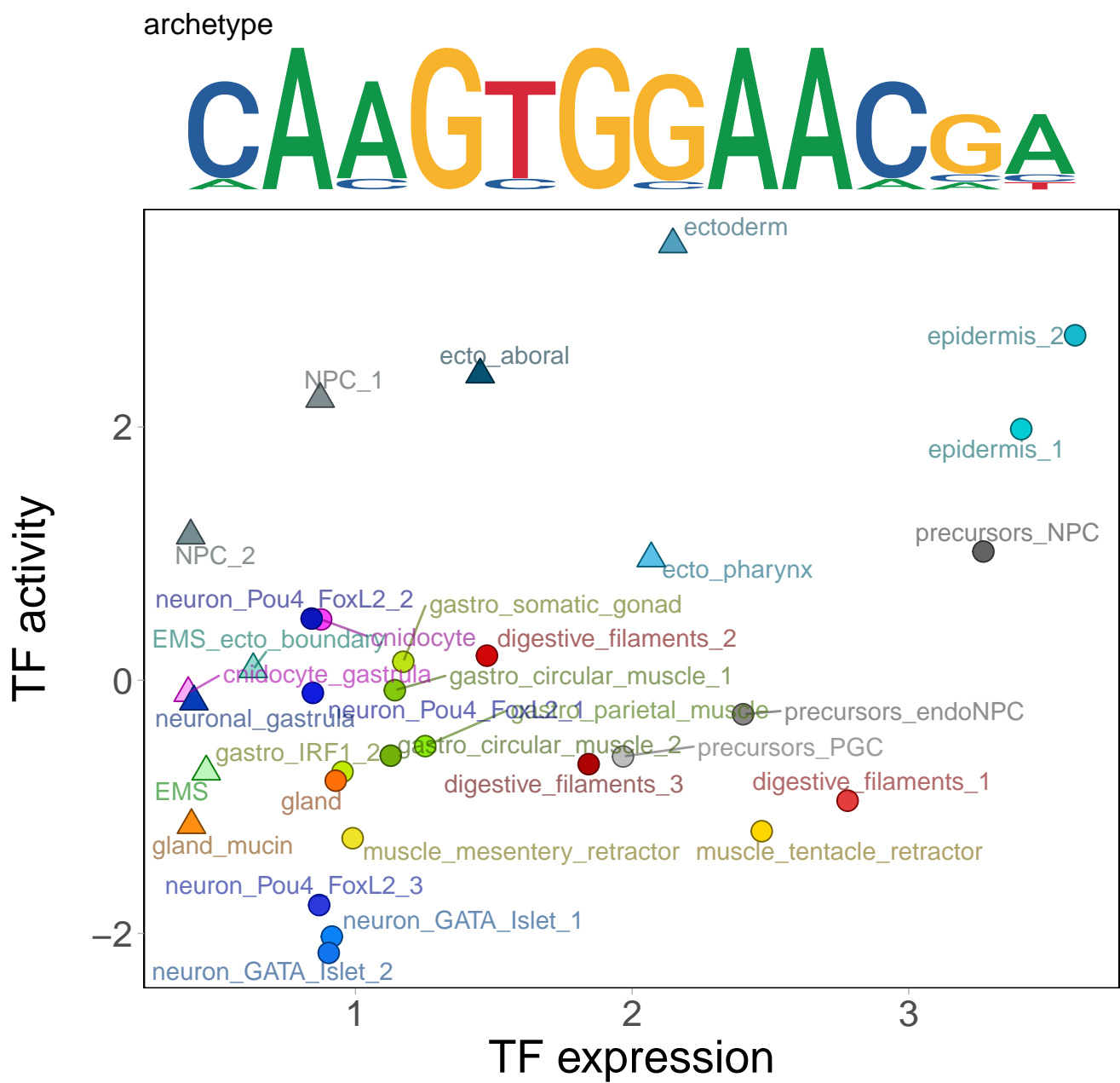
nvec_dap_5
Nvec_vc1.1_XM_032364784.2 (Hhex)
OG: Homeodomains.HG1.28:HHEX
PFAM: Homeodomain



nvec_dap_6
Nvec_vc1.1_XM_001632929.3
OG: Homeodomains.HG2.41:SHOX/SHOX2
PFAM: Homeodomain

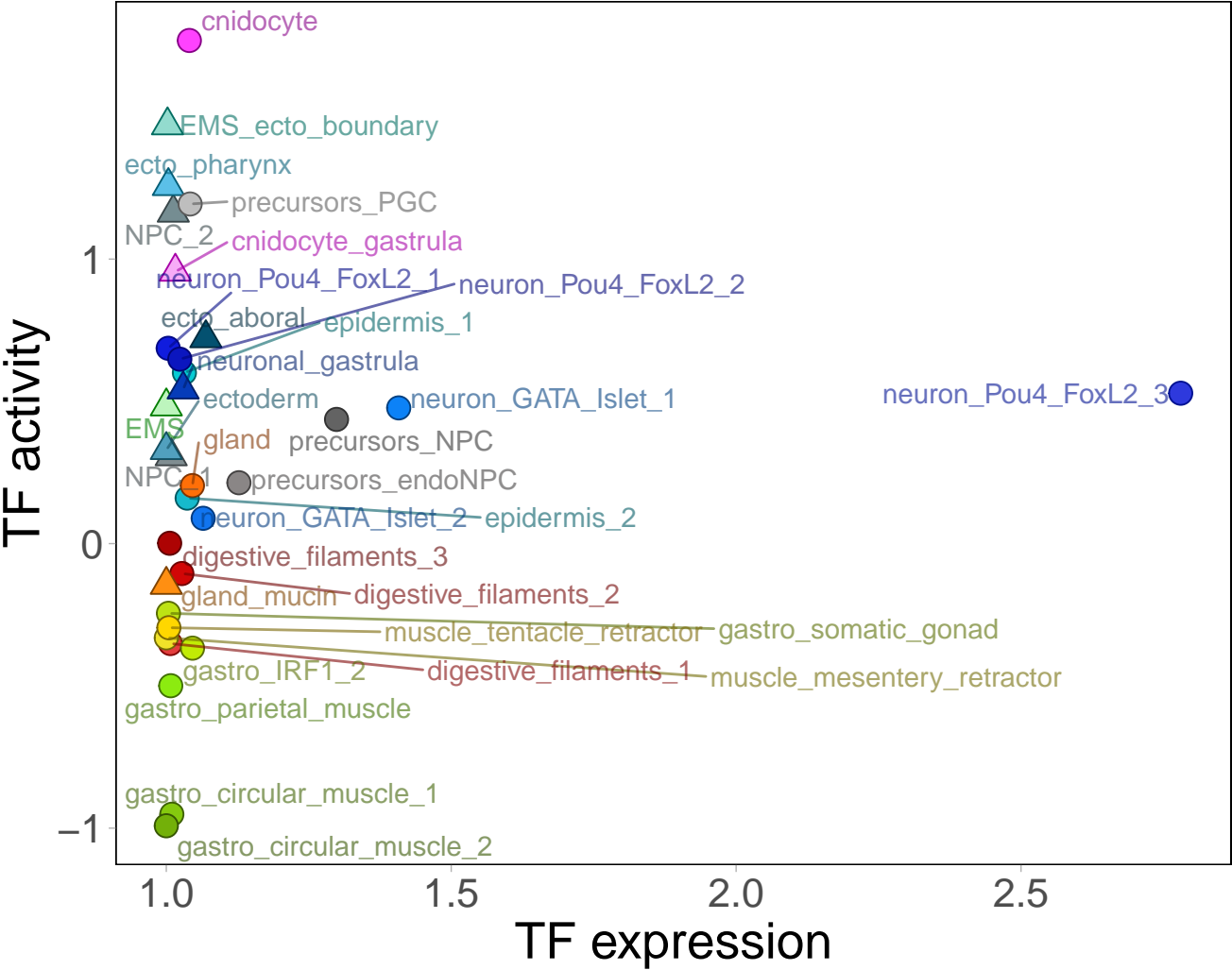


nvec_dap_8
Nvec_vc1.1_XM_032384889.2 (Hes2)
OG: HLH.HG1.63:like:HES5/HES7
PFAM: HLH/Hairy_orange

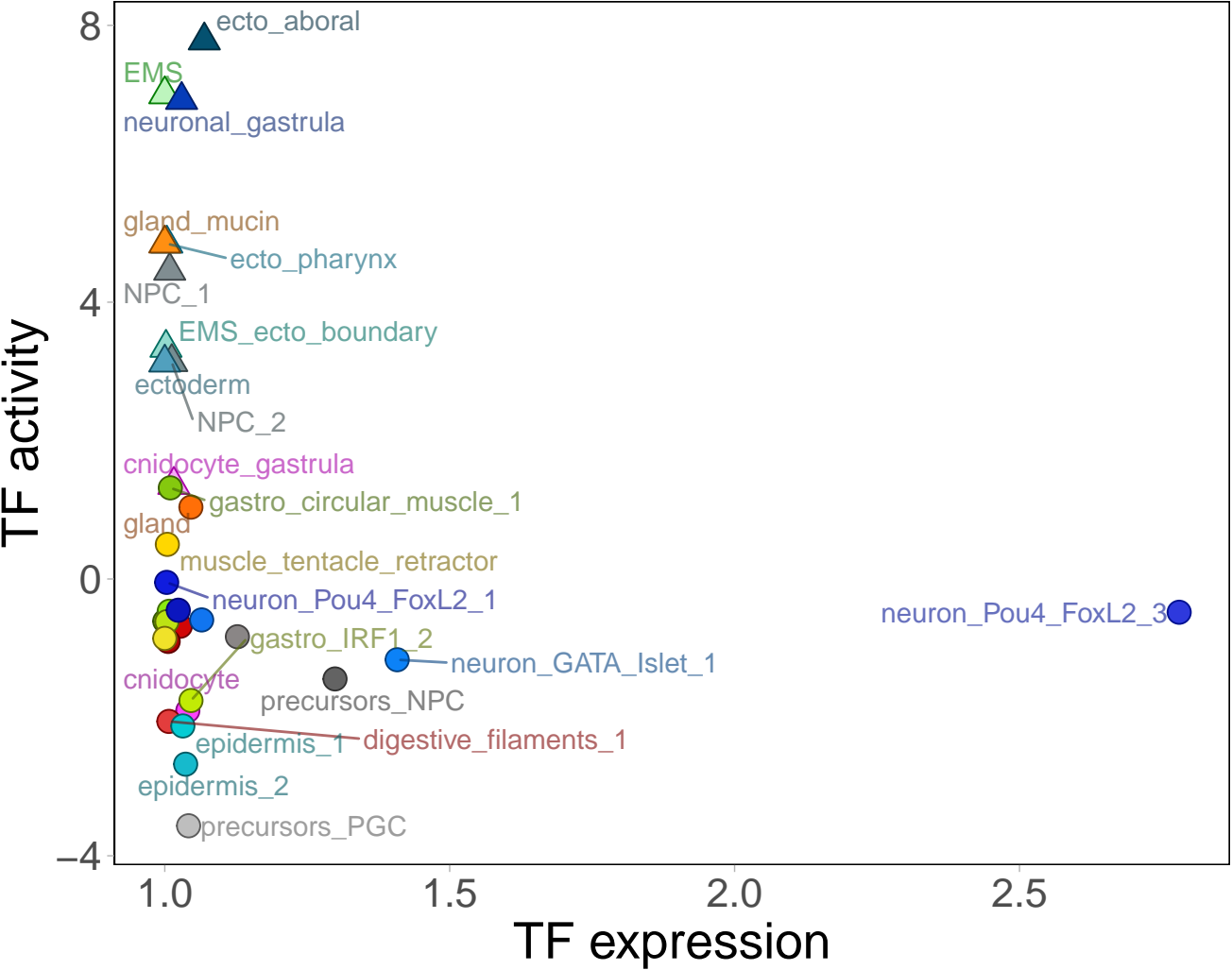


nvec_dap_10
Nvec_vc1.1_XM_001626717.3
OG: T-box.HG1.4:like:EOMES/TBR1/TBX19/TBX21/TBXT
PFAM: T-box

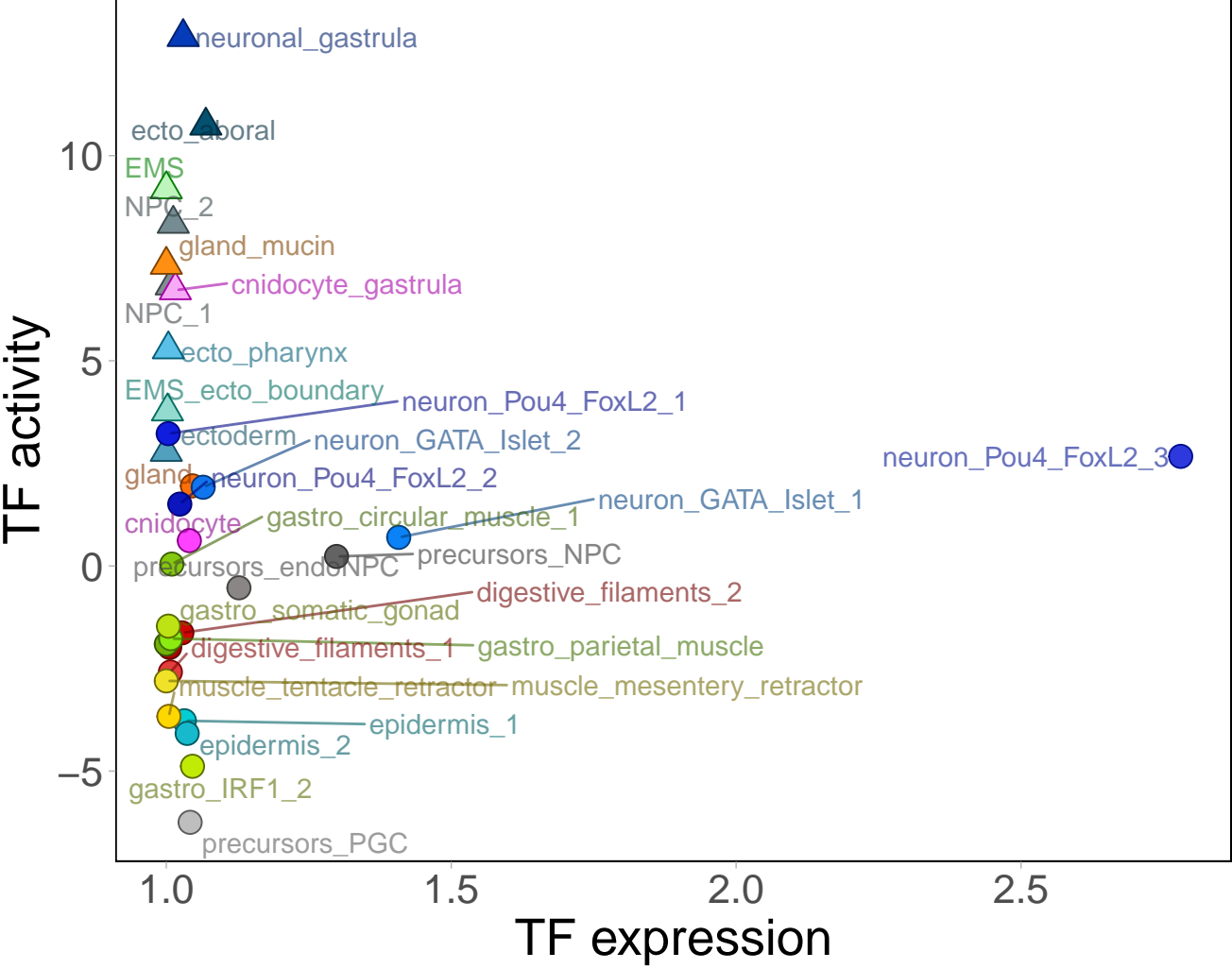
archetype



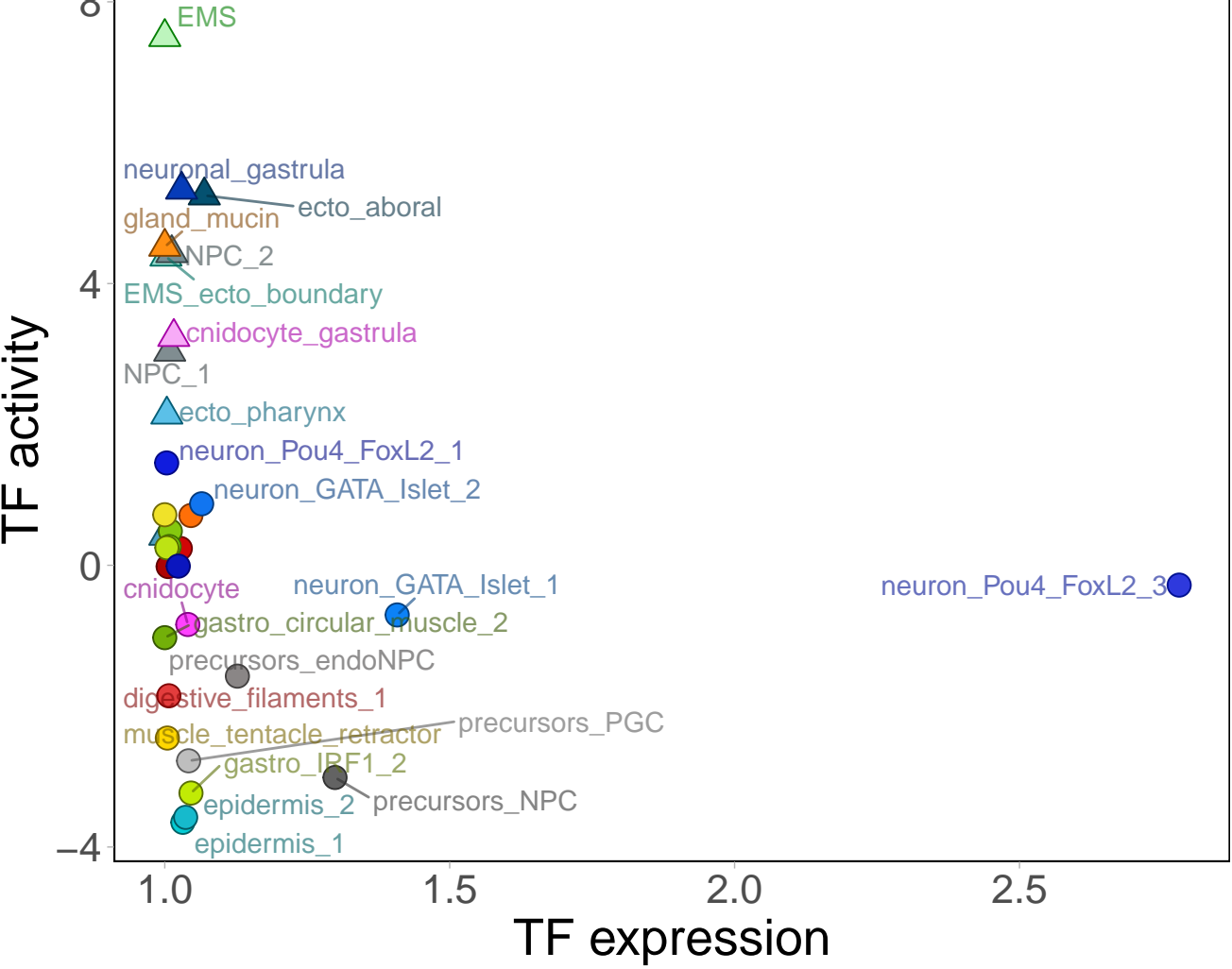
dimont



homer

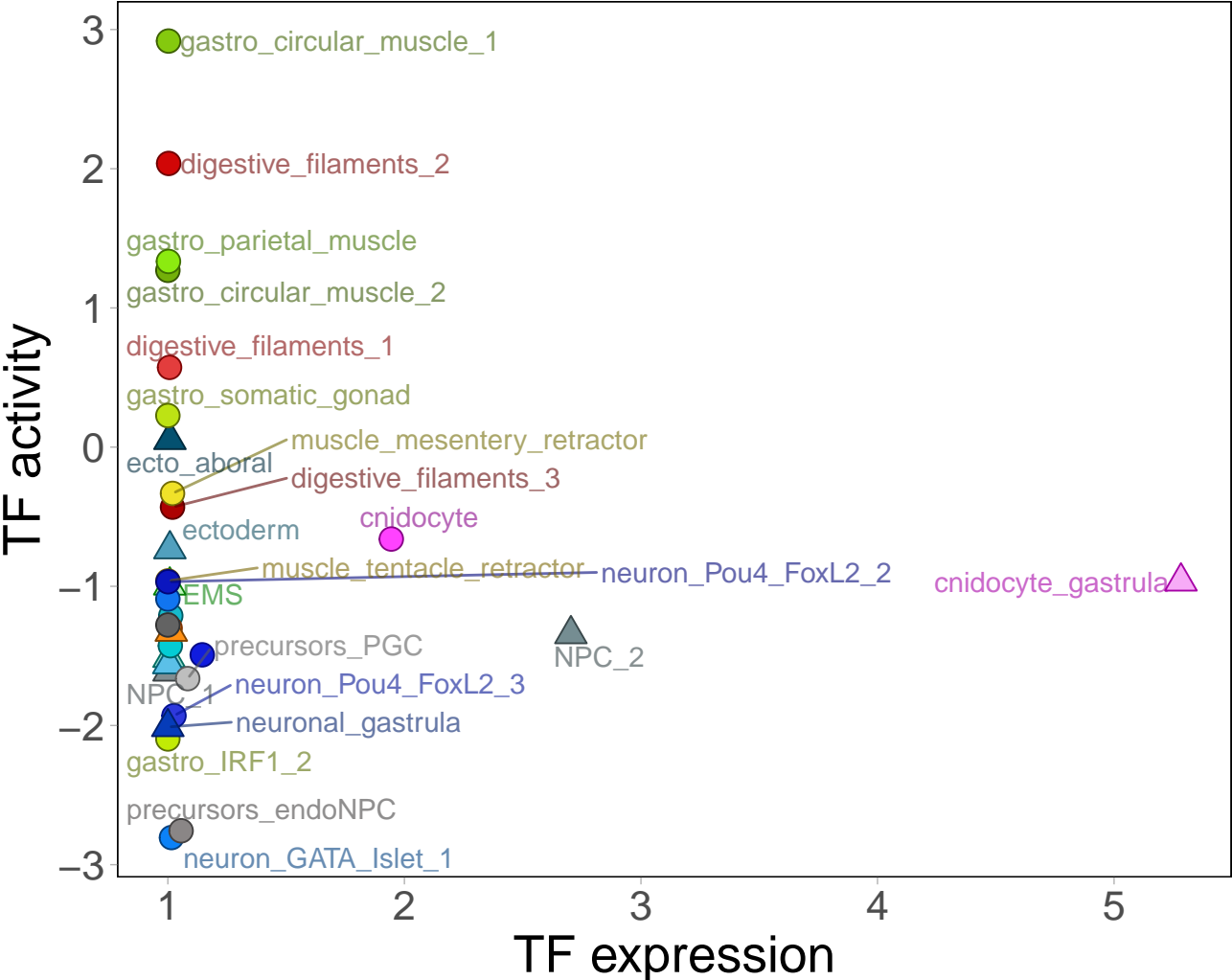


streame

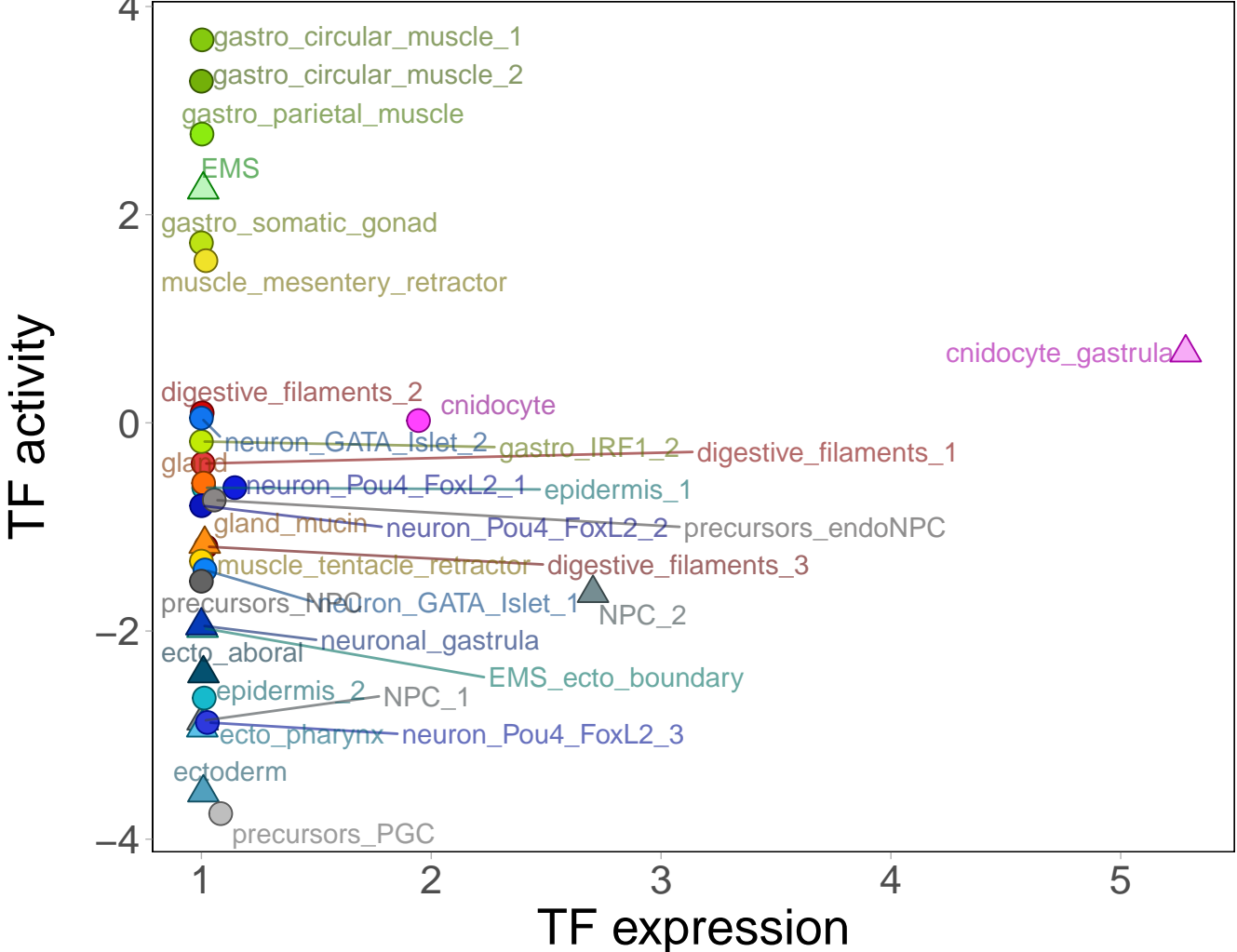


nvec_dap_11
Nvec_vc1.1_XM_032364331.2
OG: TEA.HG1.0:TEAD1/TEAD2/TEAD3/TEAD4
PFAM: TEA

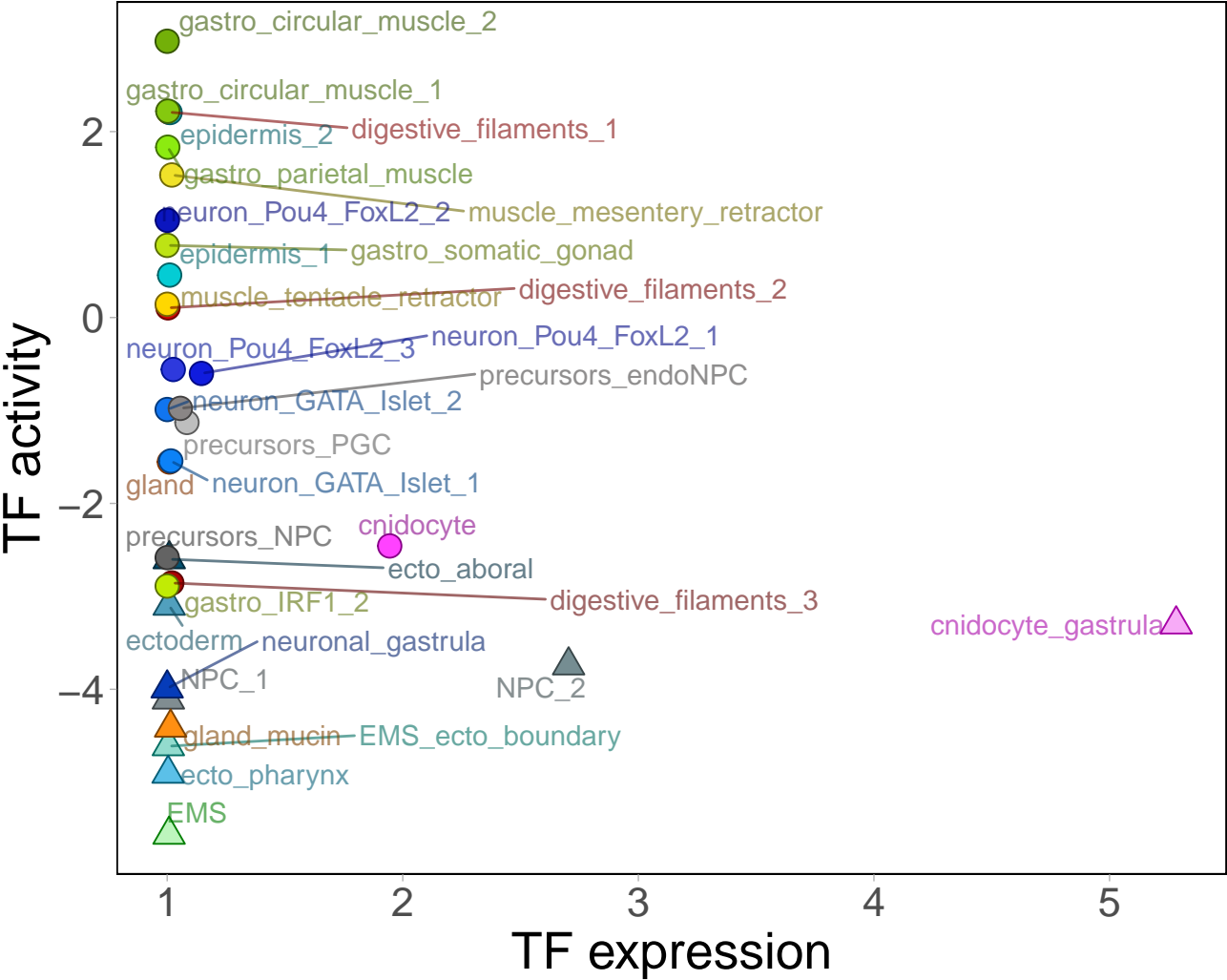
archetype



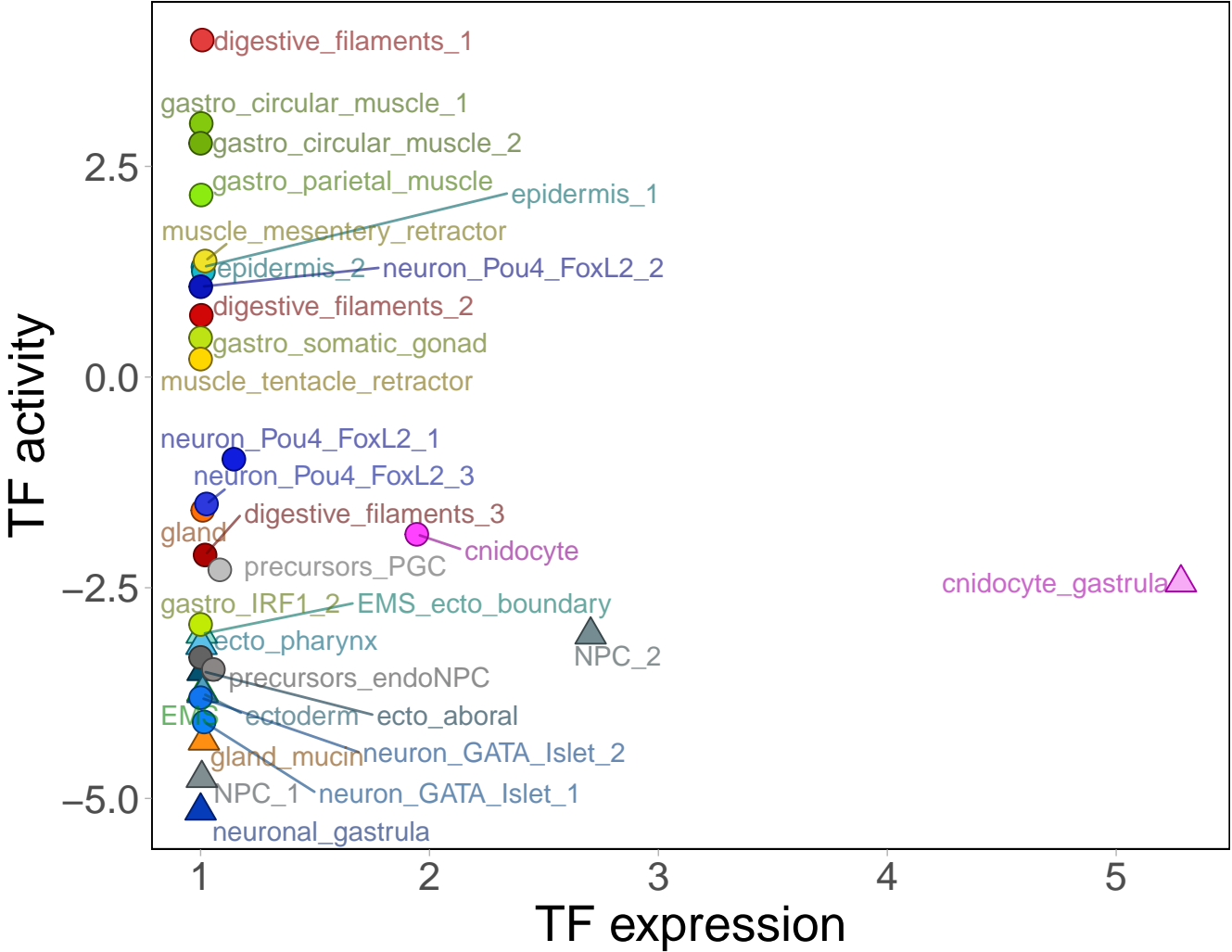
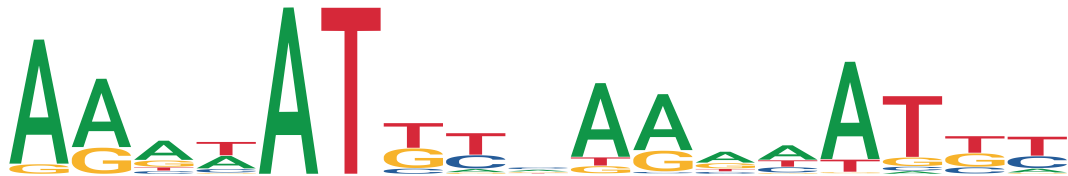
dimont



homer

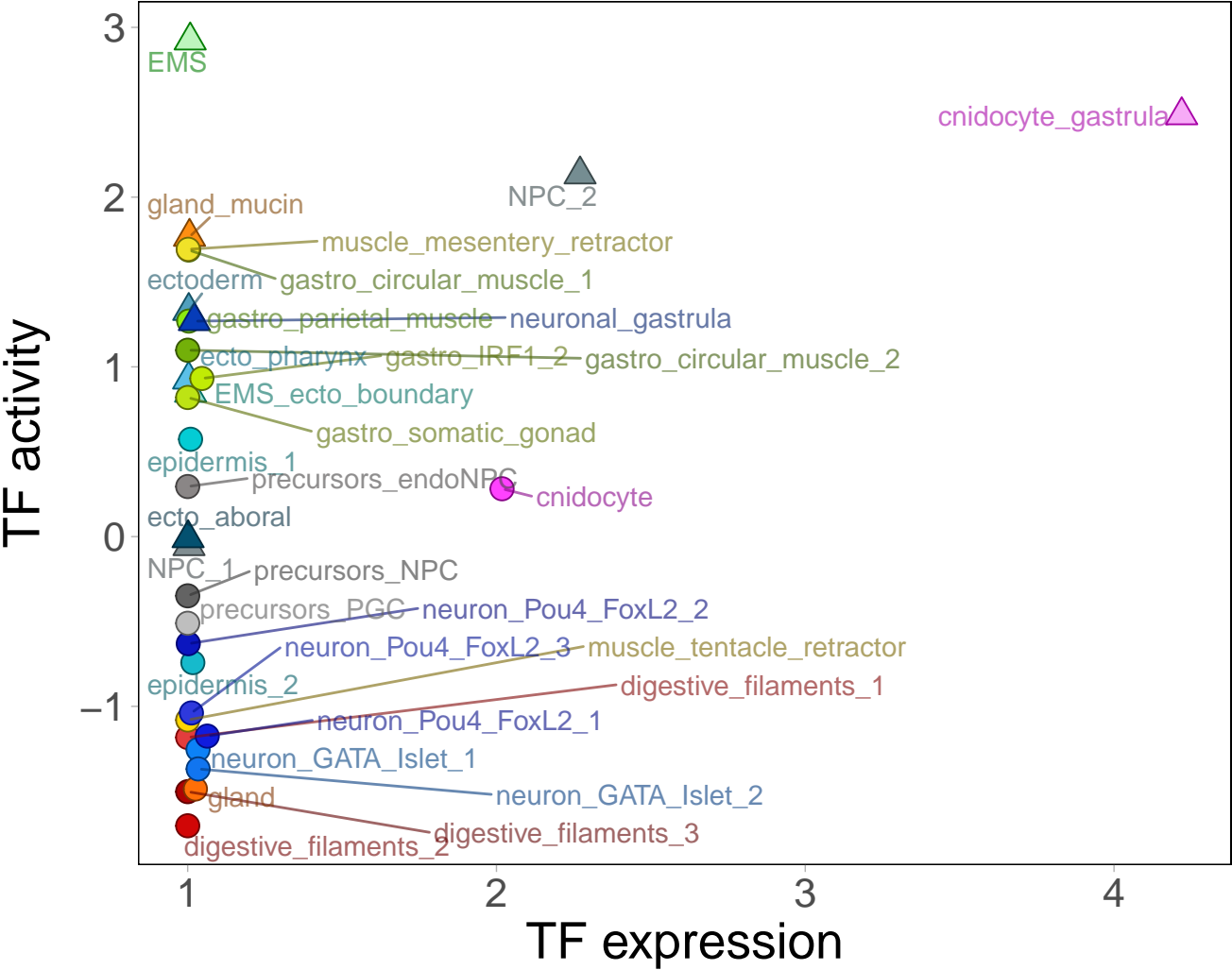


streame

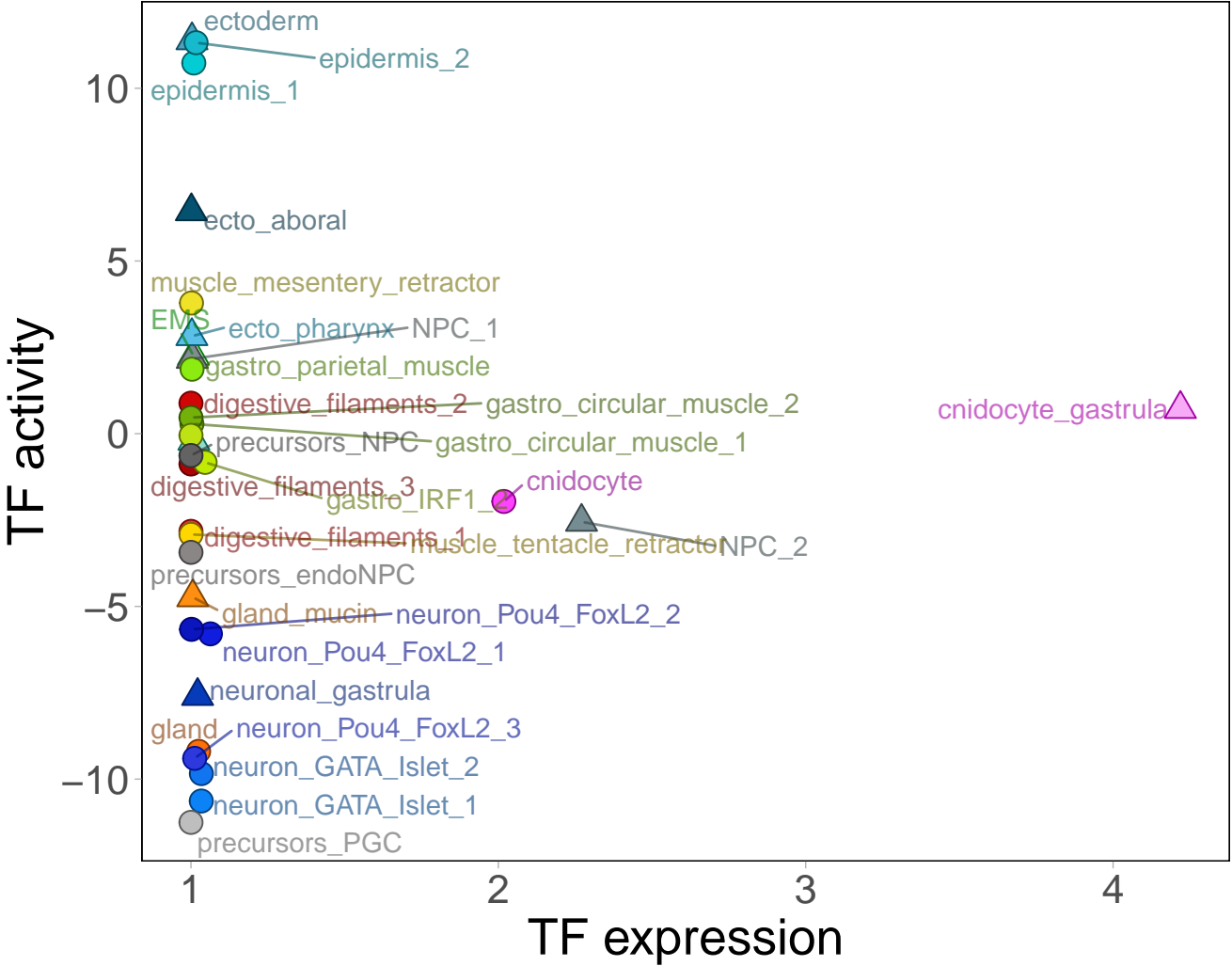


nvec_dap_13
Nvec_vc1.1_XM_001631763.3 (PaxA)
OG: Homeodomains.HG2.49:like:PAX1/PAX9
PFAM: PAX

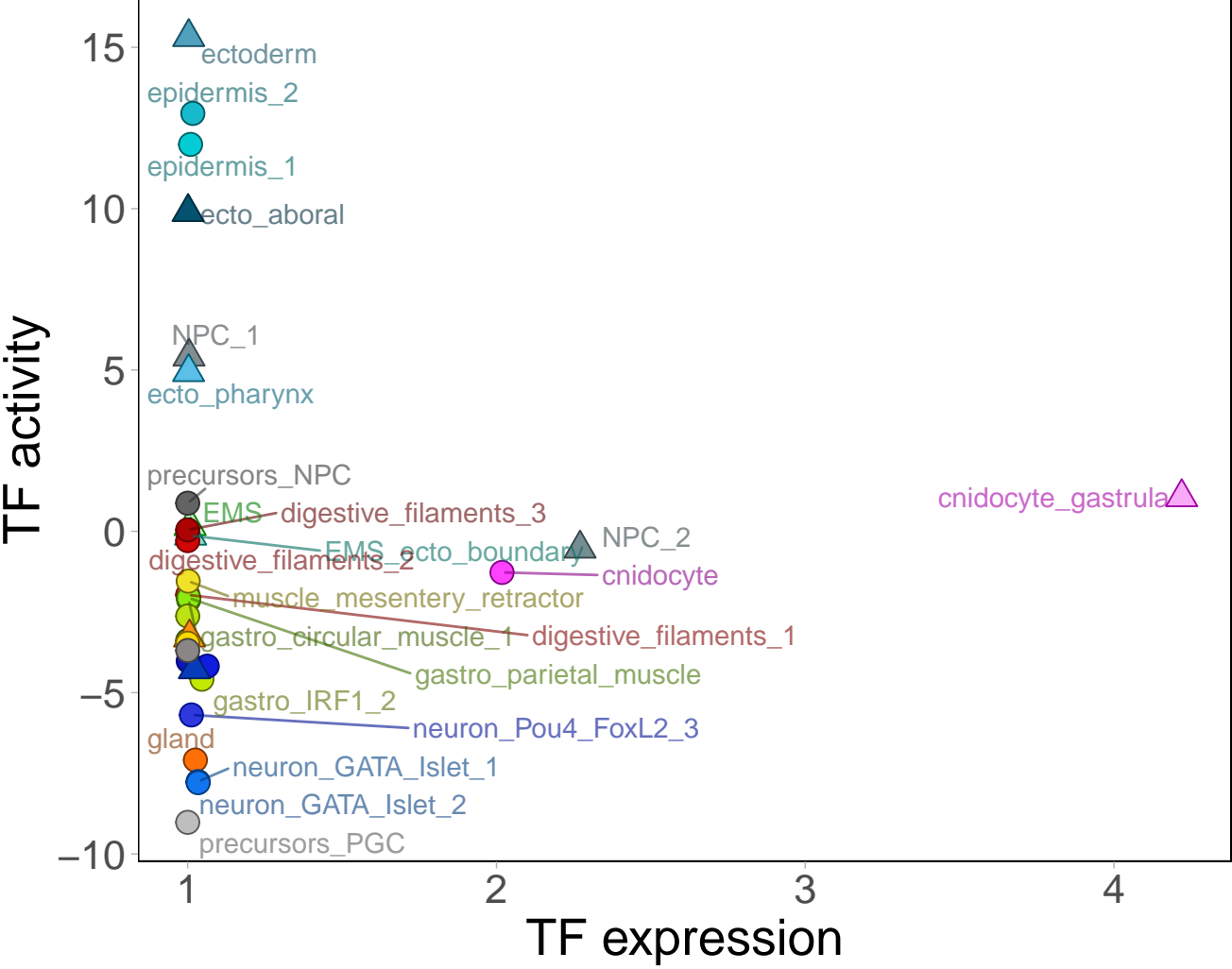
archetype



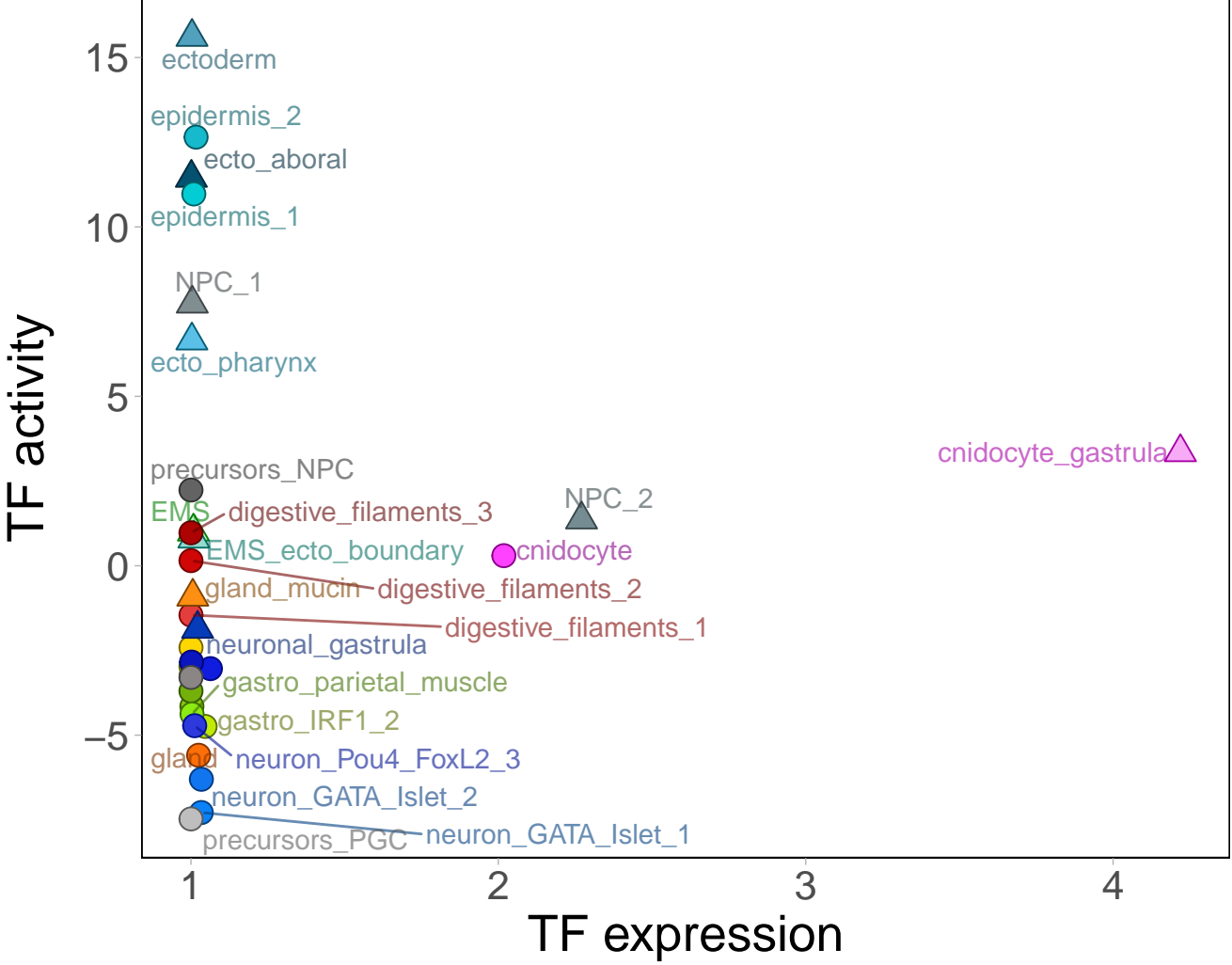
dimont



homer

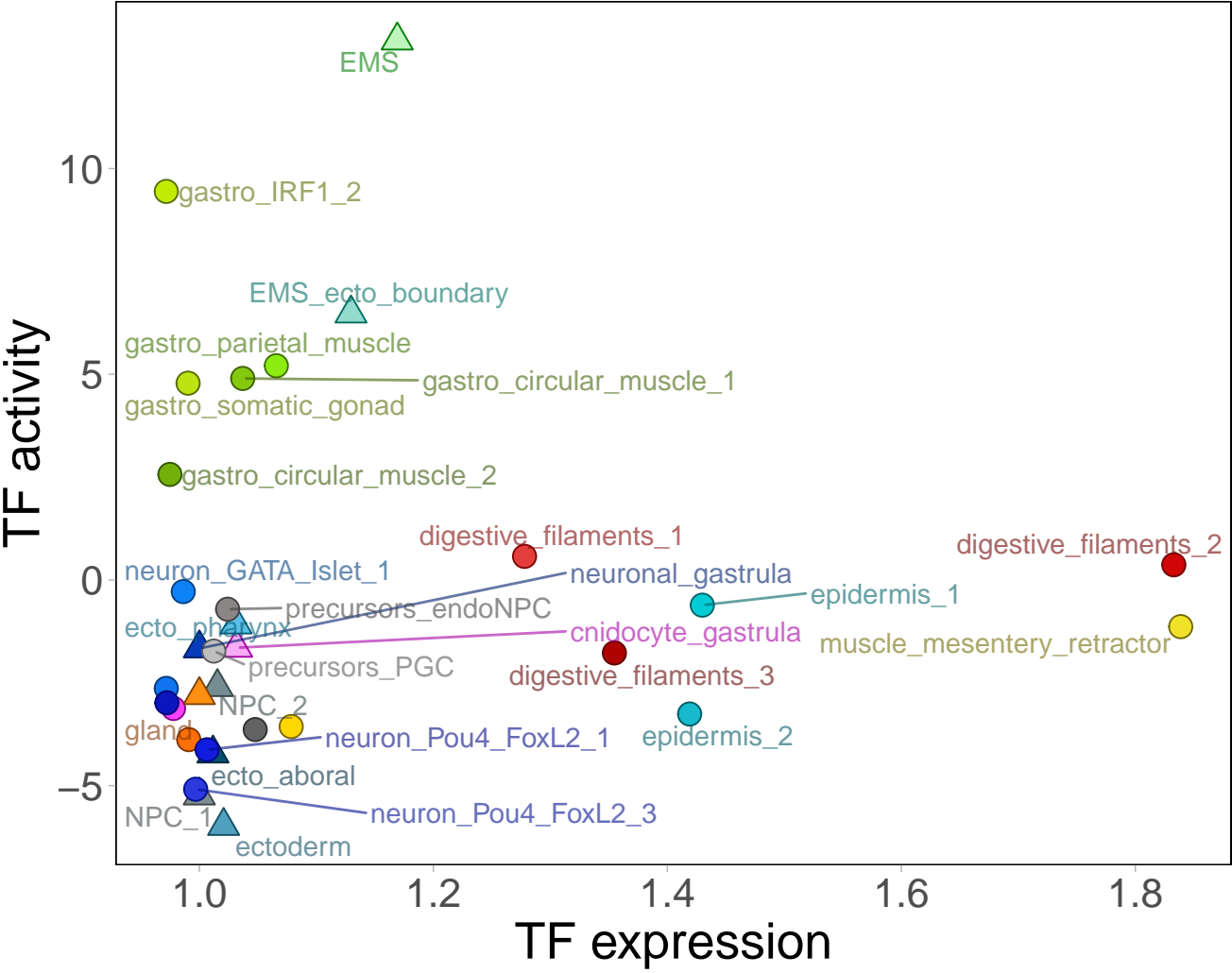


streme

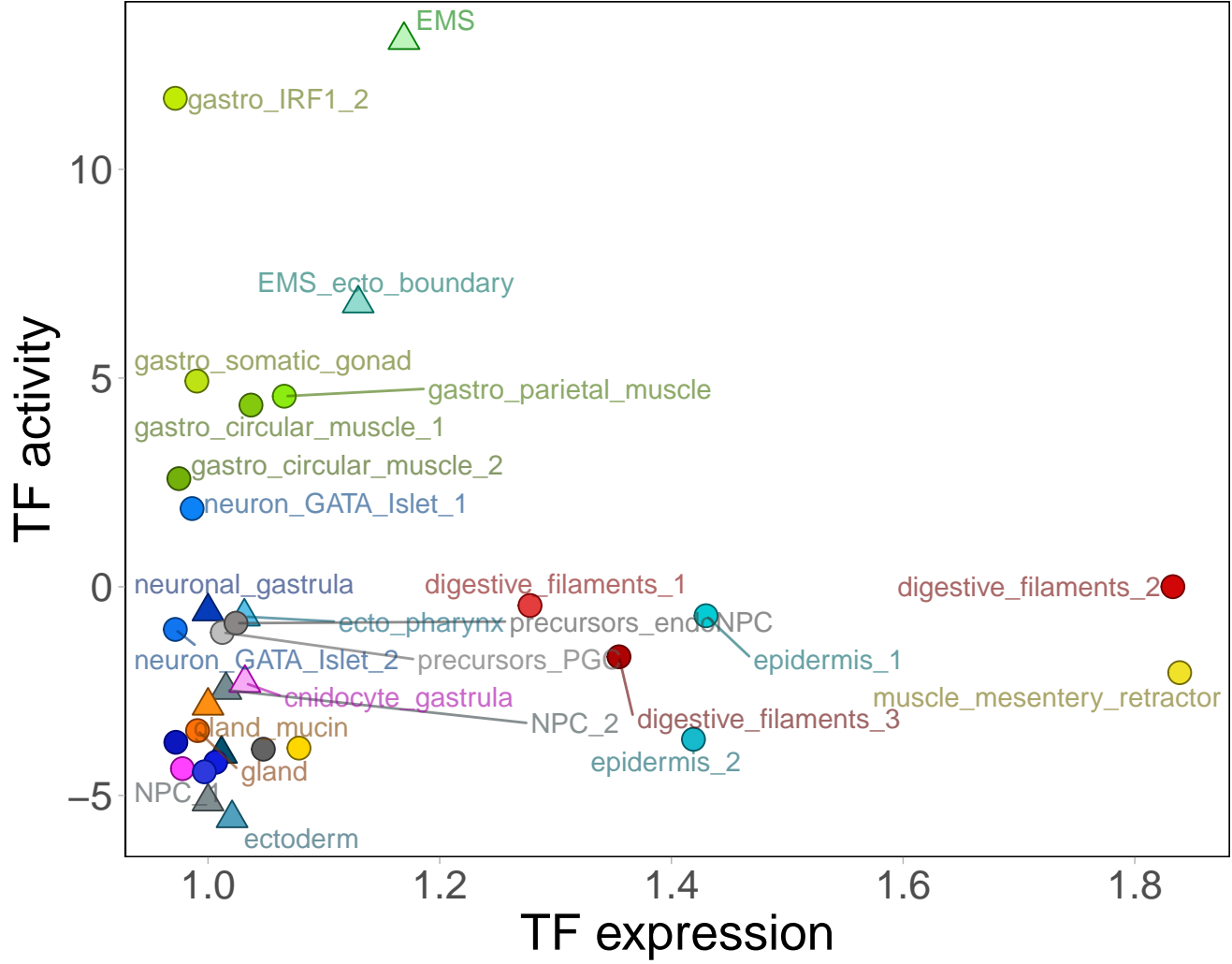


nvec_dap_14
Nvec_vc1.1_XM_001637374.3
OG: Ets.HG1.2:ELK1/ELK3/ELK4
PFAM: Ets

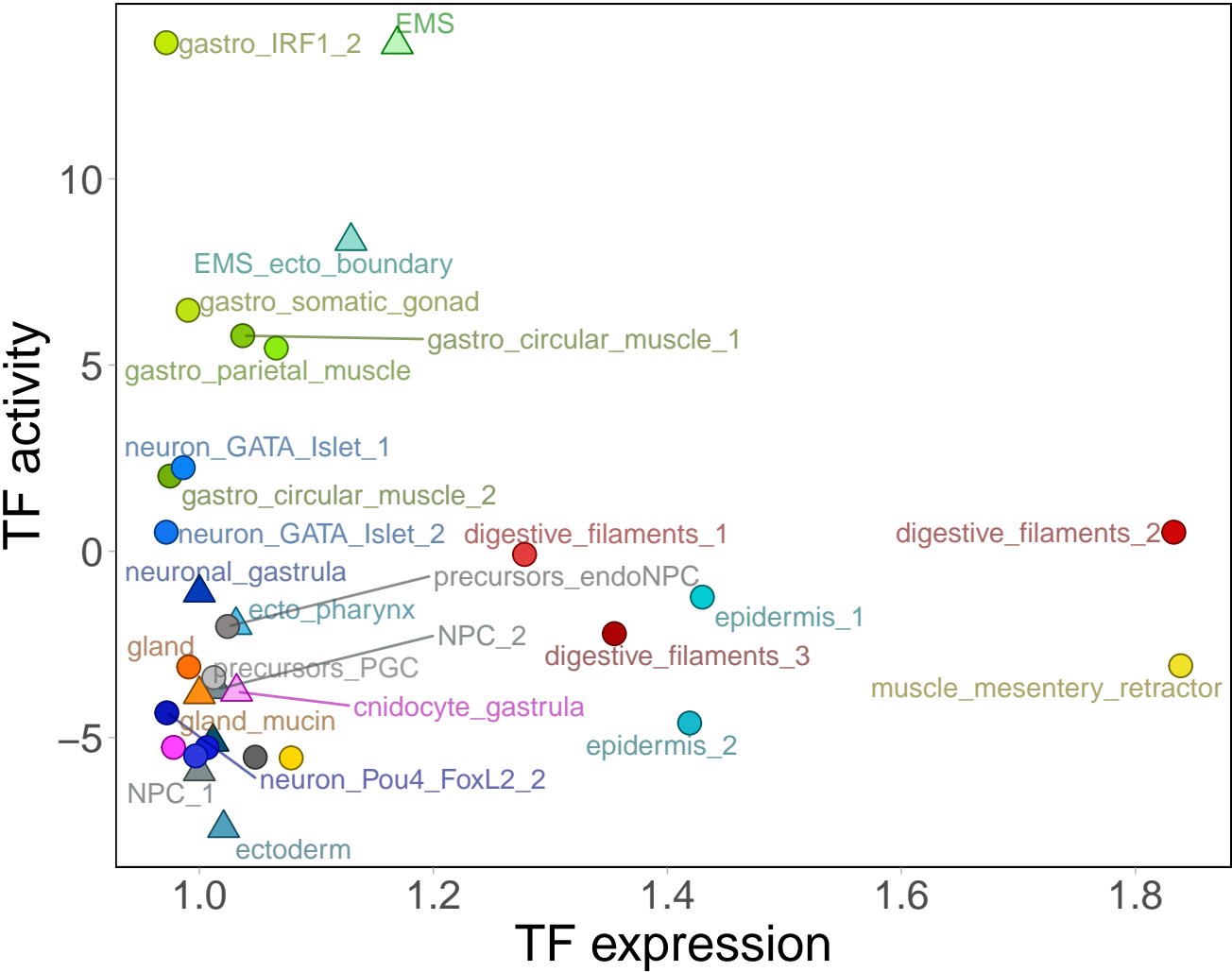
archetype



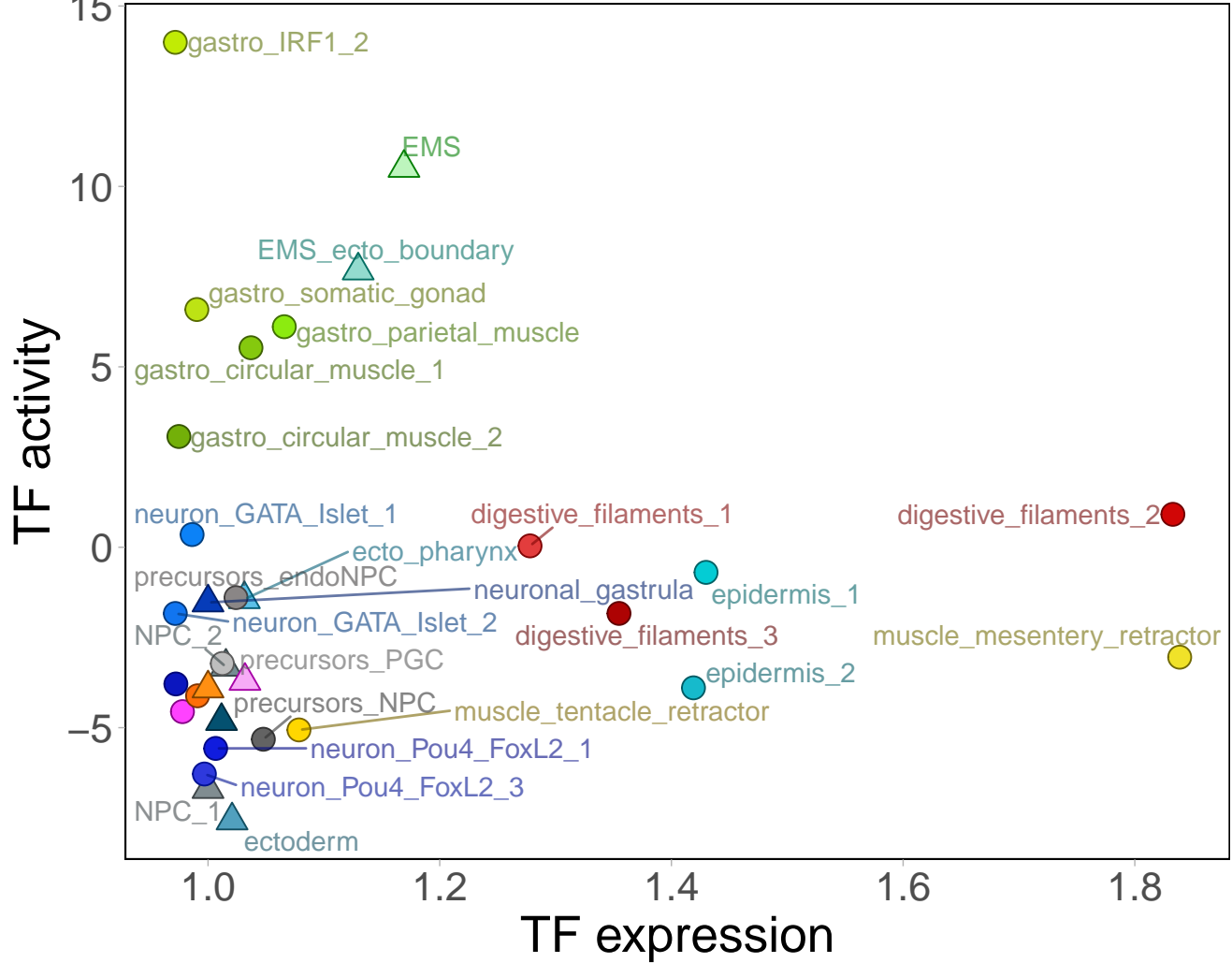
dimont



homer

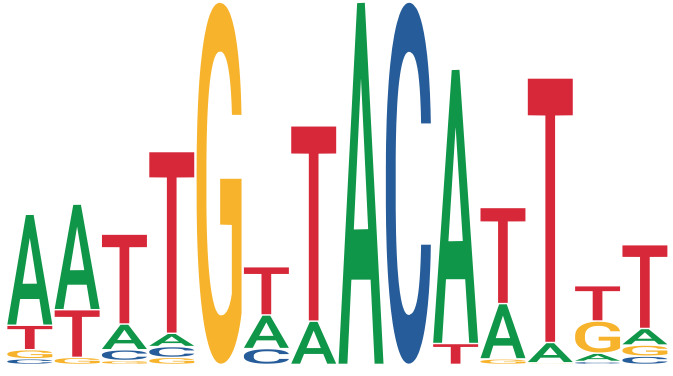


streame



nvec_dap_15
Nvec_vc1.1_XM_032371744.2
OG: Doublesex_DM.HG1.12:DMRTA1/DMRTA2/DMRTB1
PFAM: DM/DMA

dimont



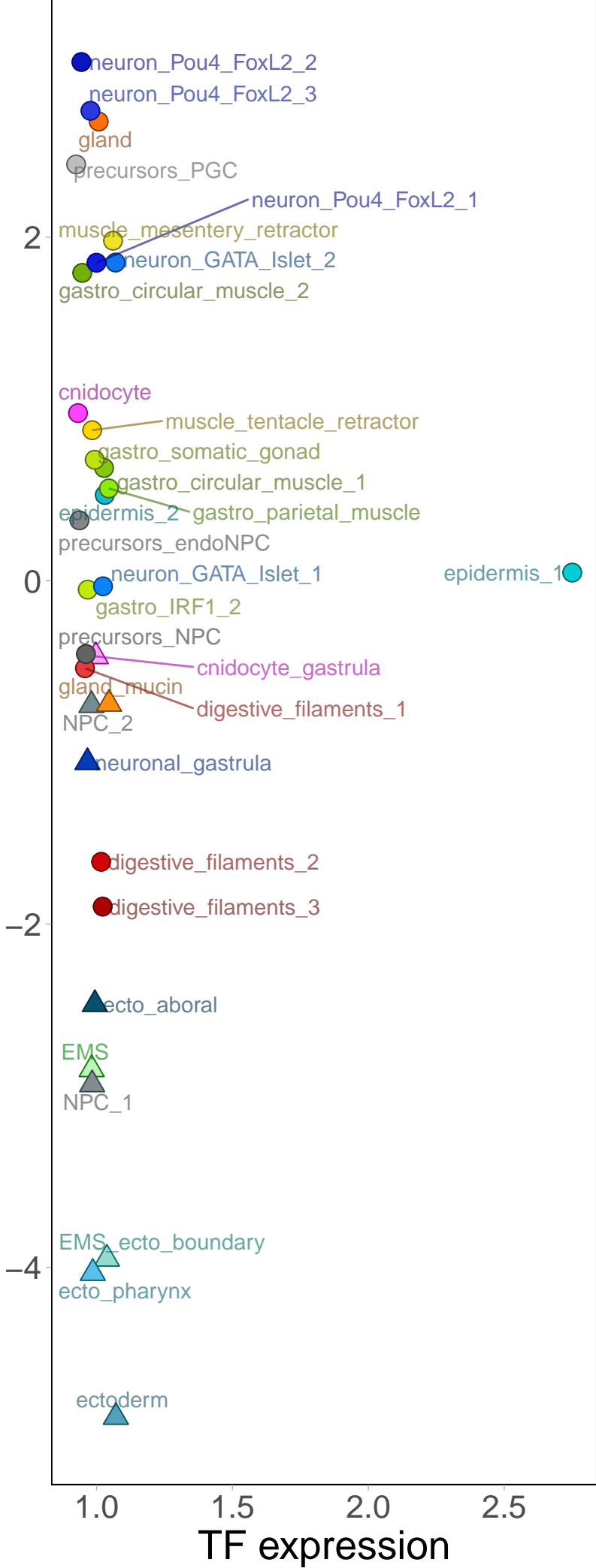
homer



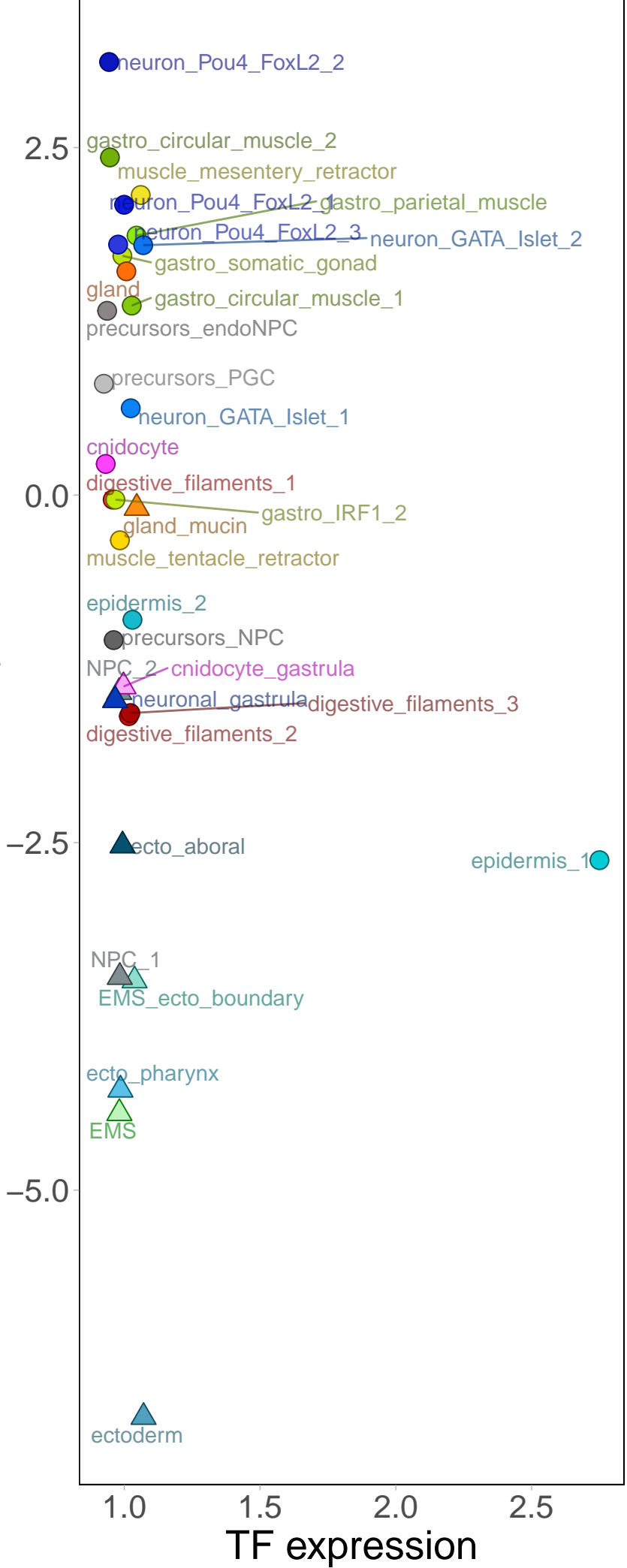
streme



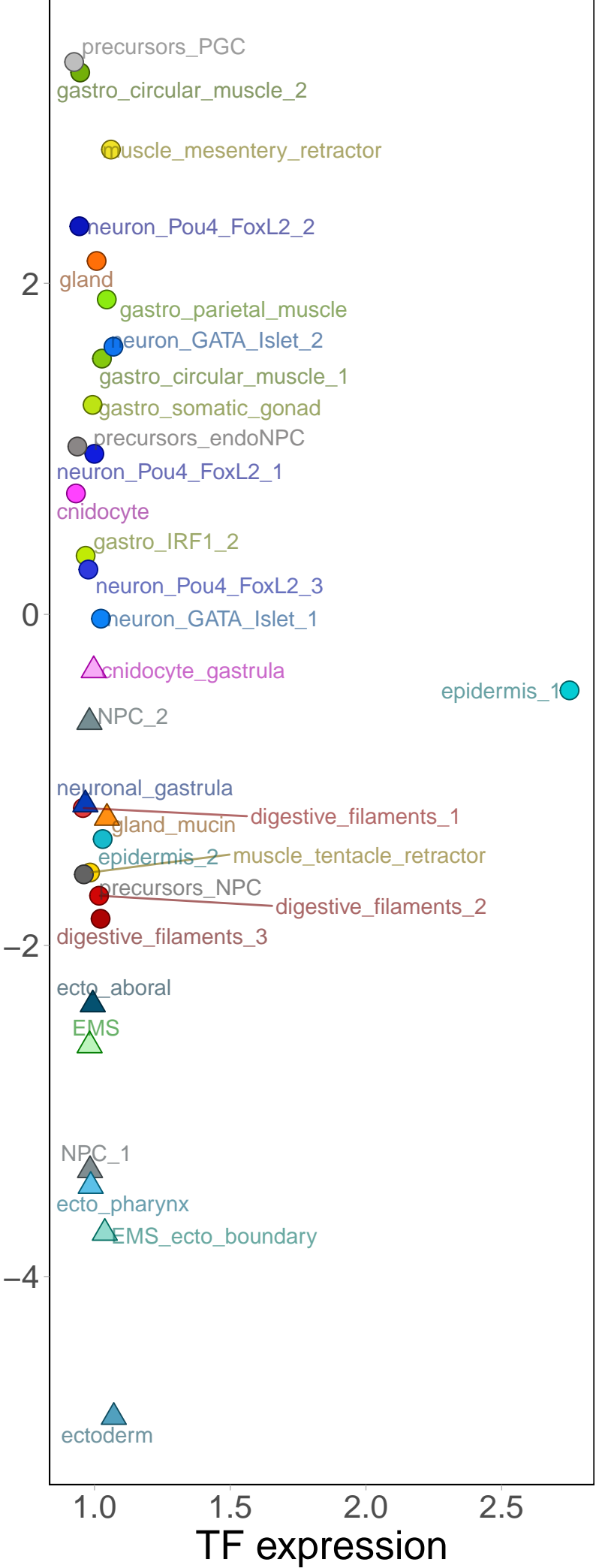
TF activity



TF activity

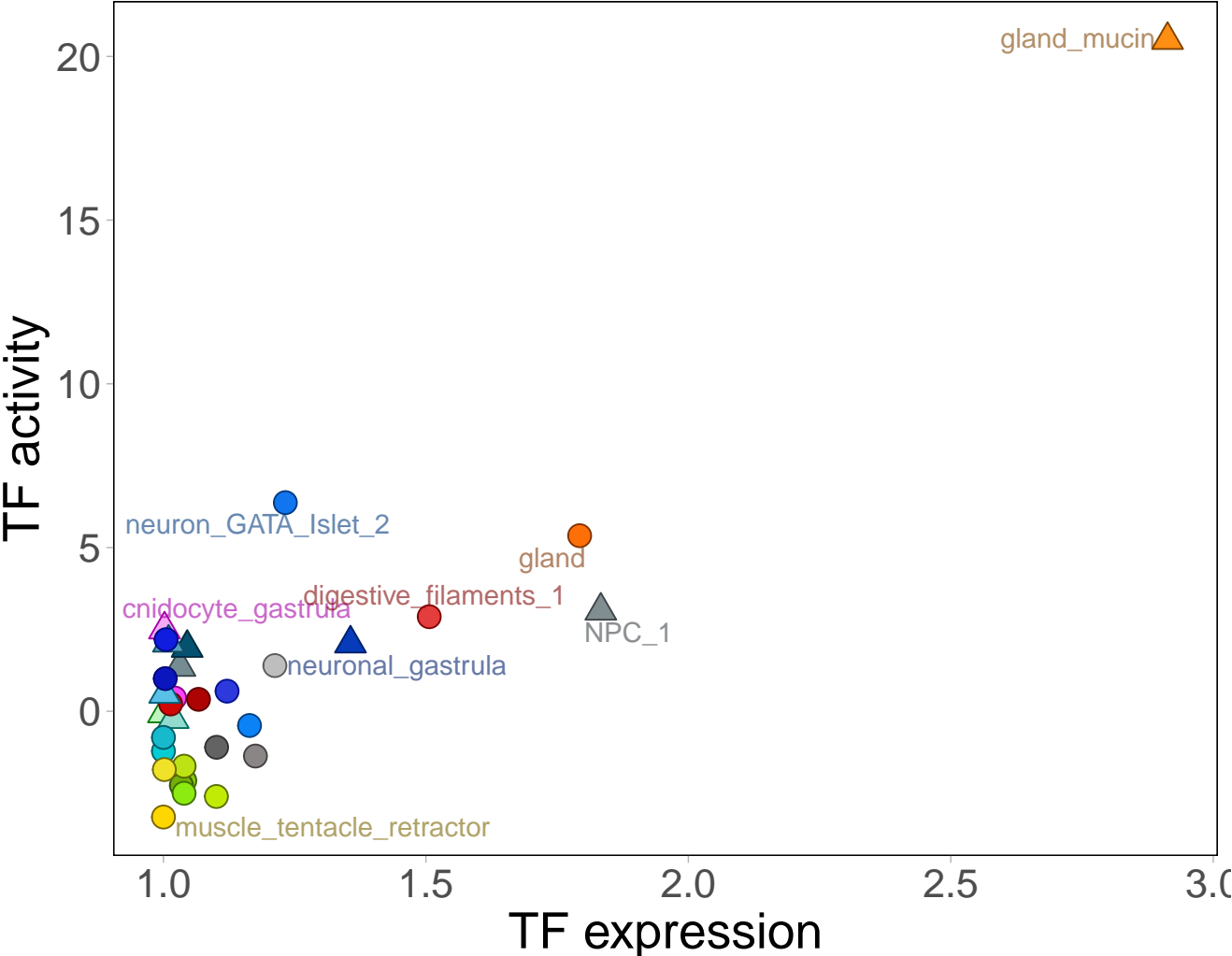


TF activity

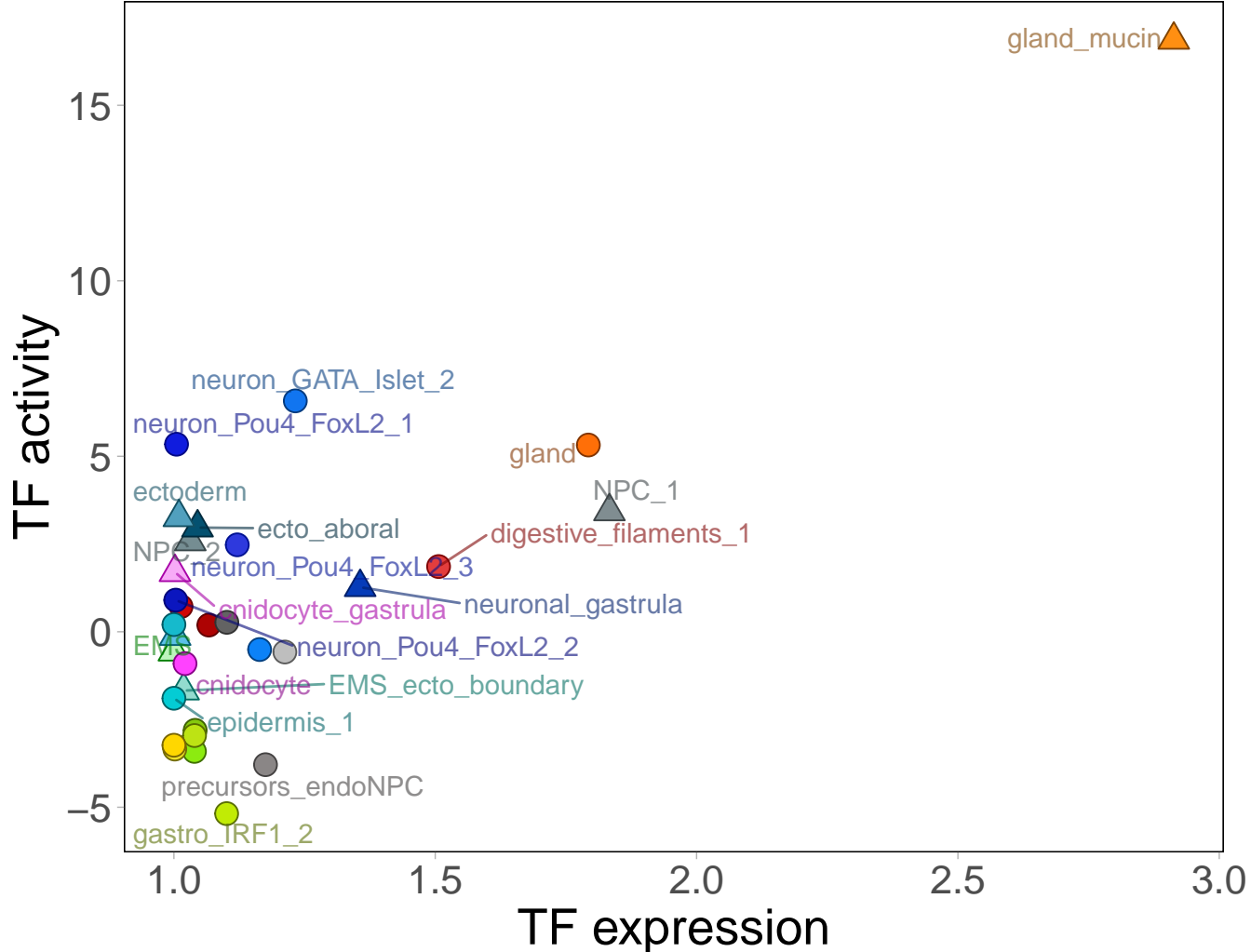


nvec_dap_16
Nvec_vc1.1_XM_032375228.2 (Rfx4)
OG: RFX.HG1.3:like:RFX4/RFX6/RFX8
PFAM: RFX_DNA_binding

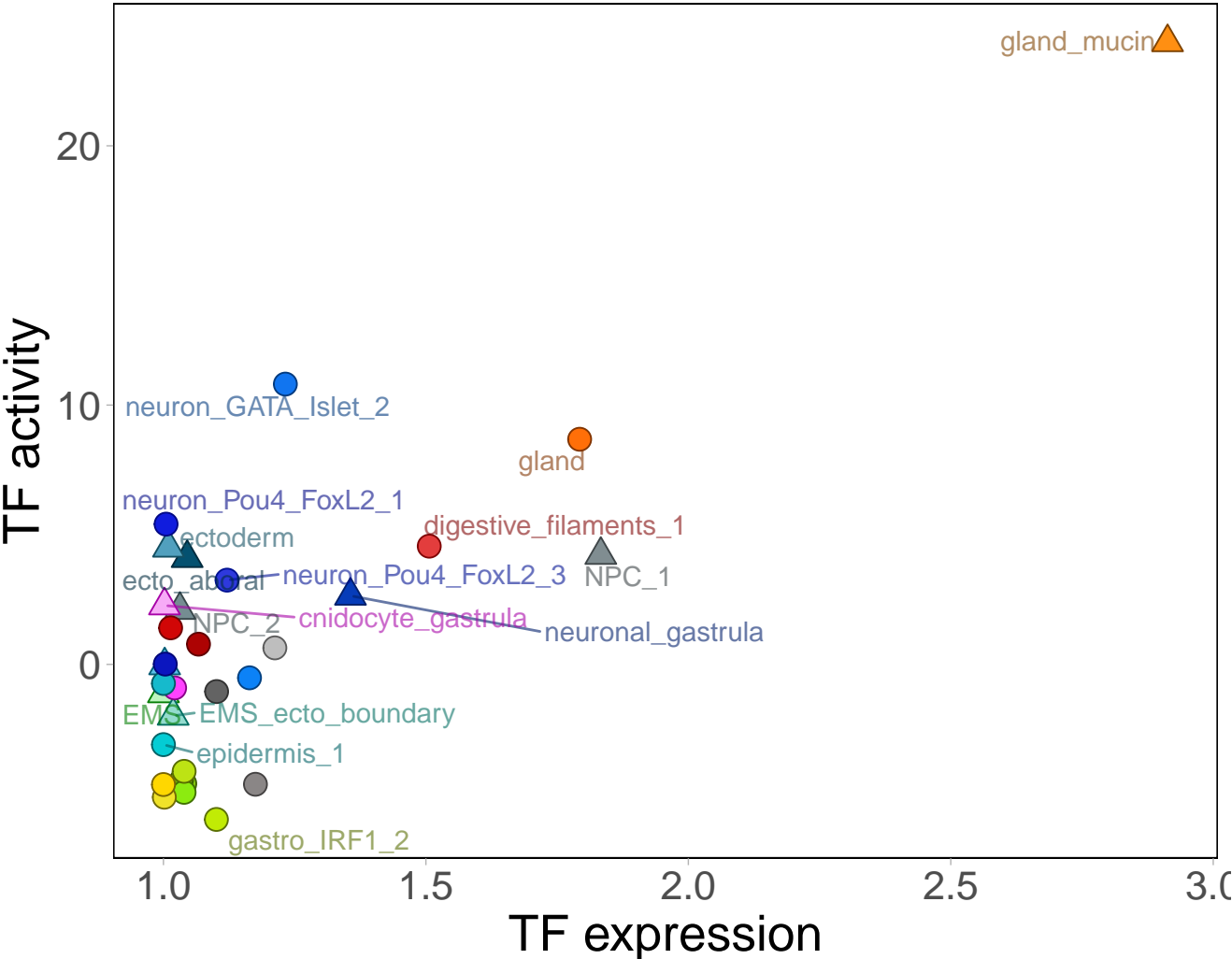
archetype



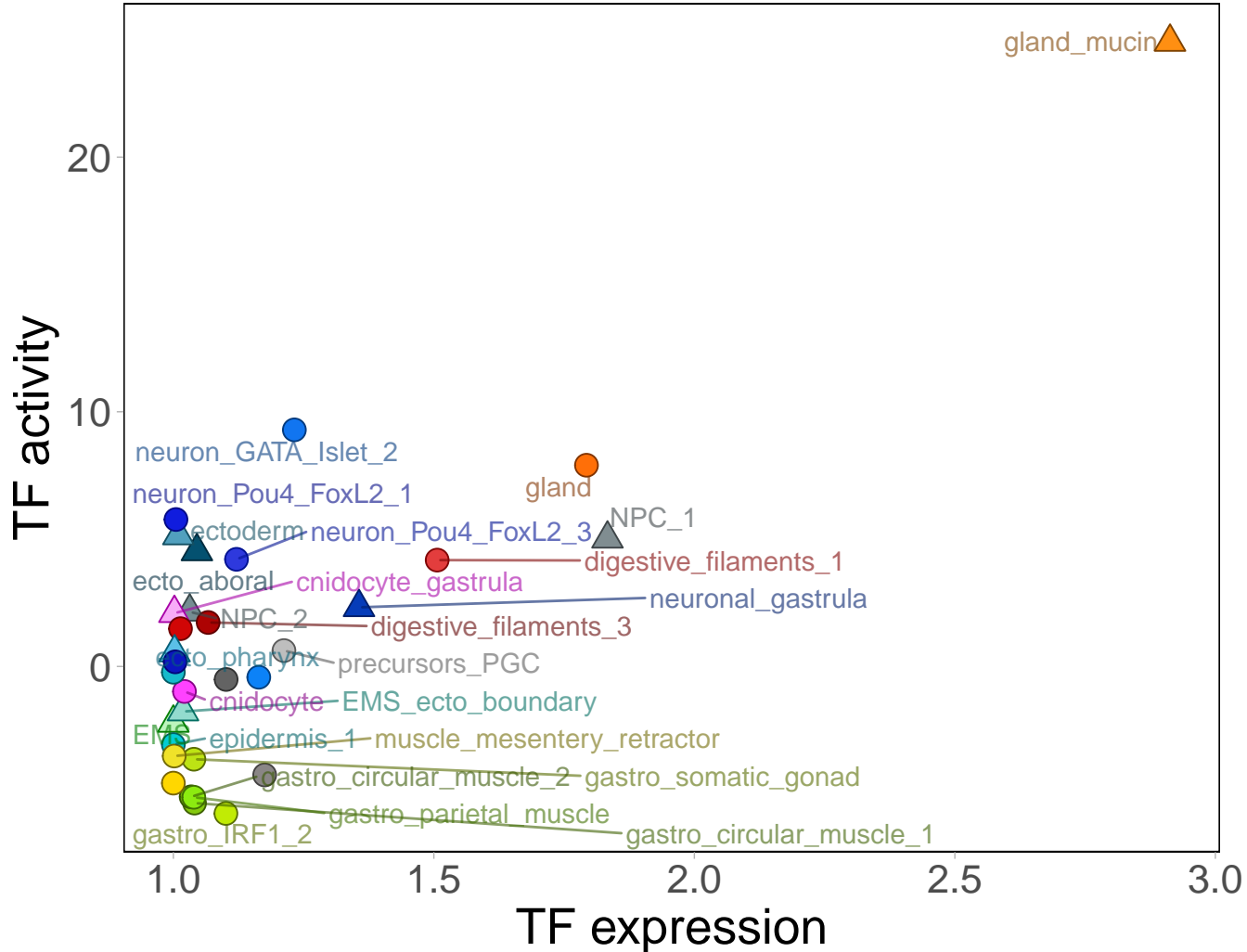
dimont



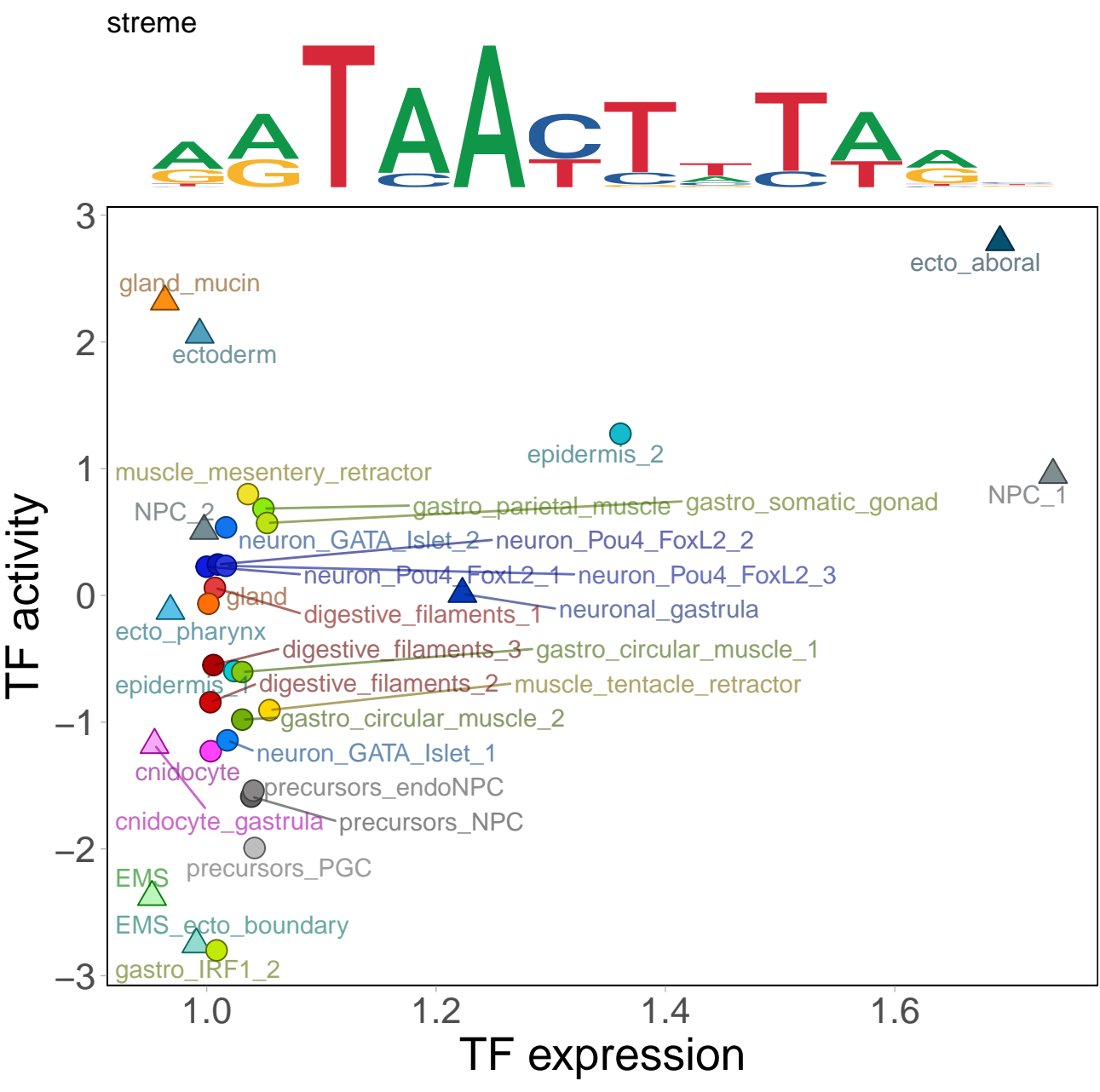
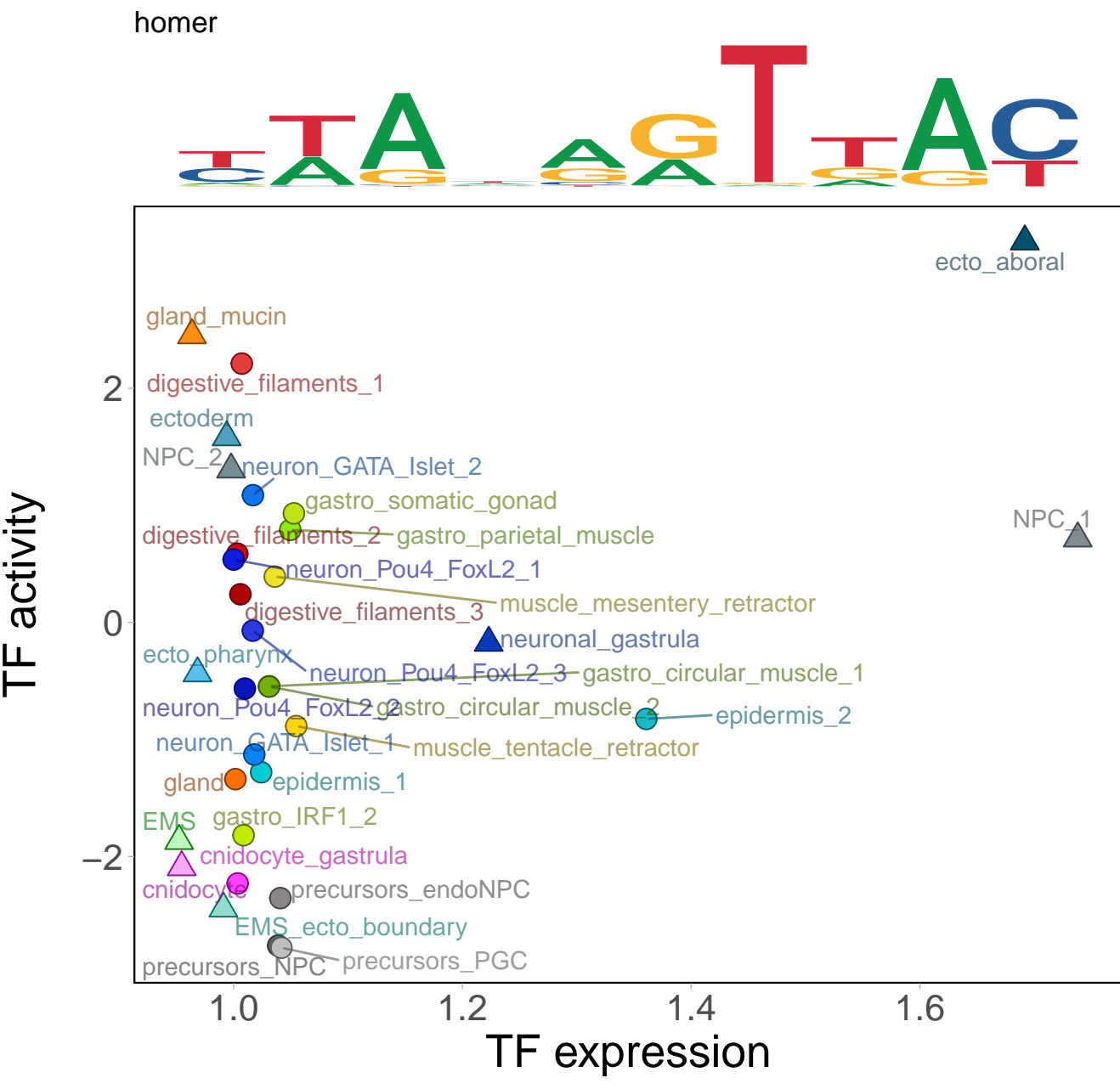
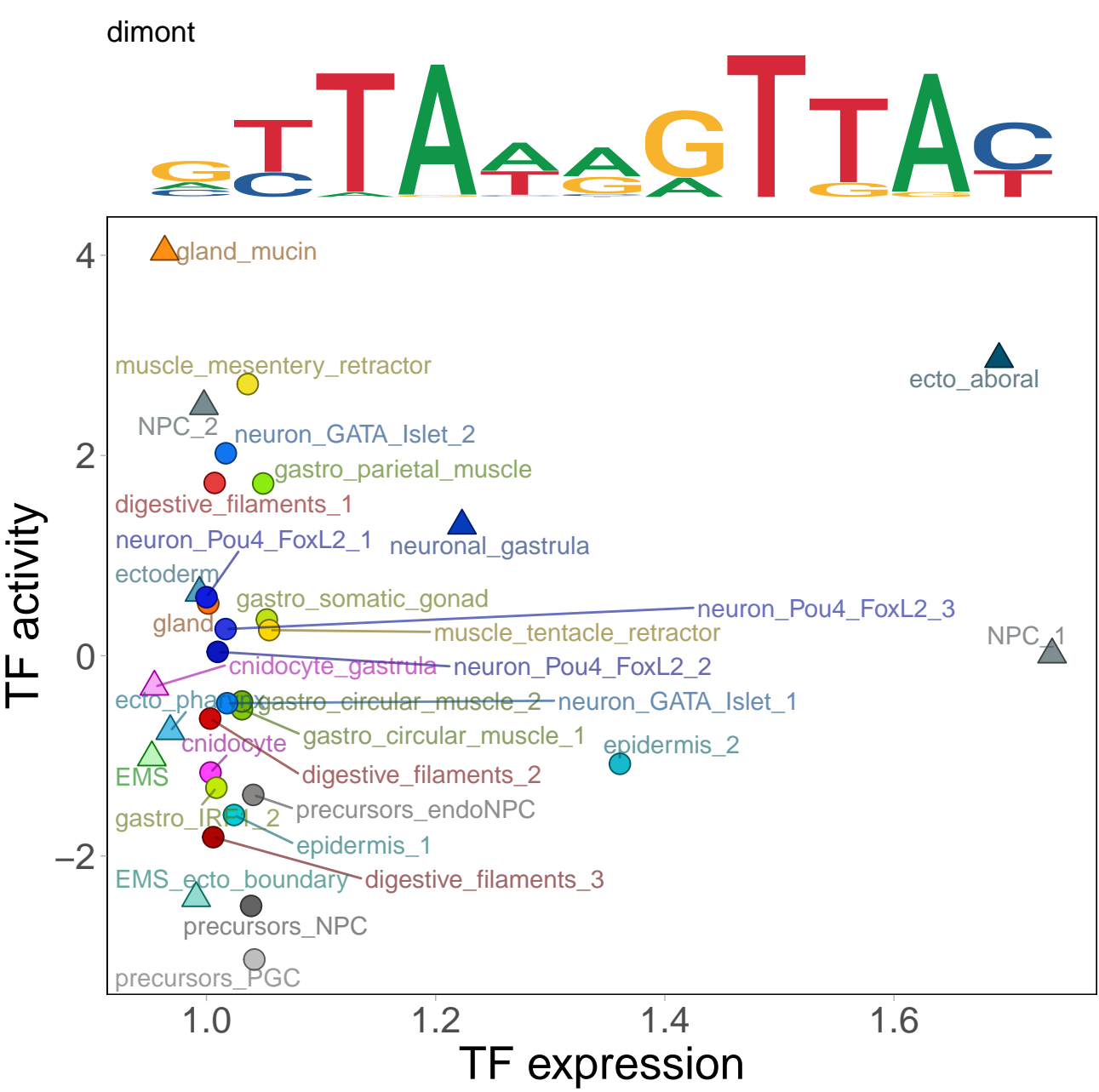
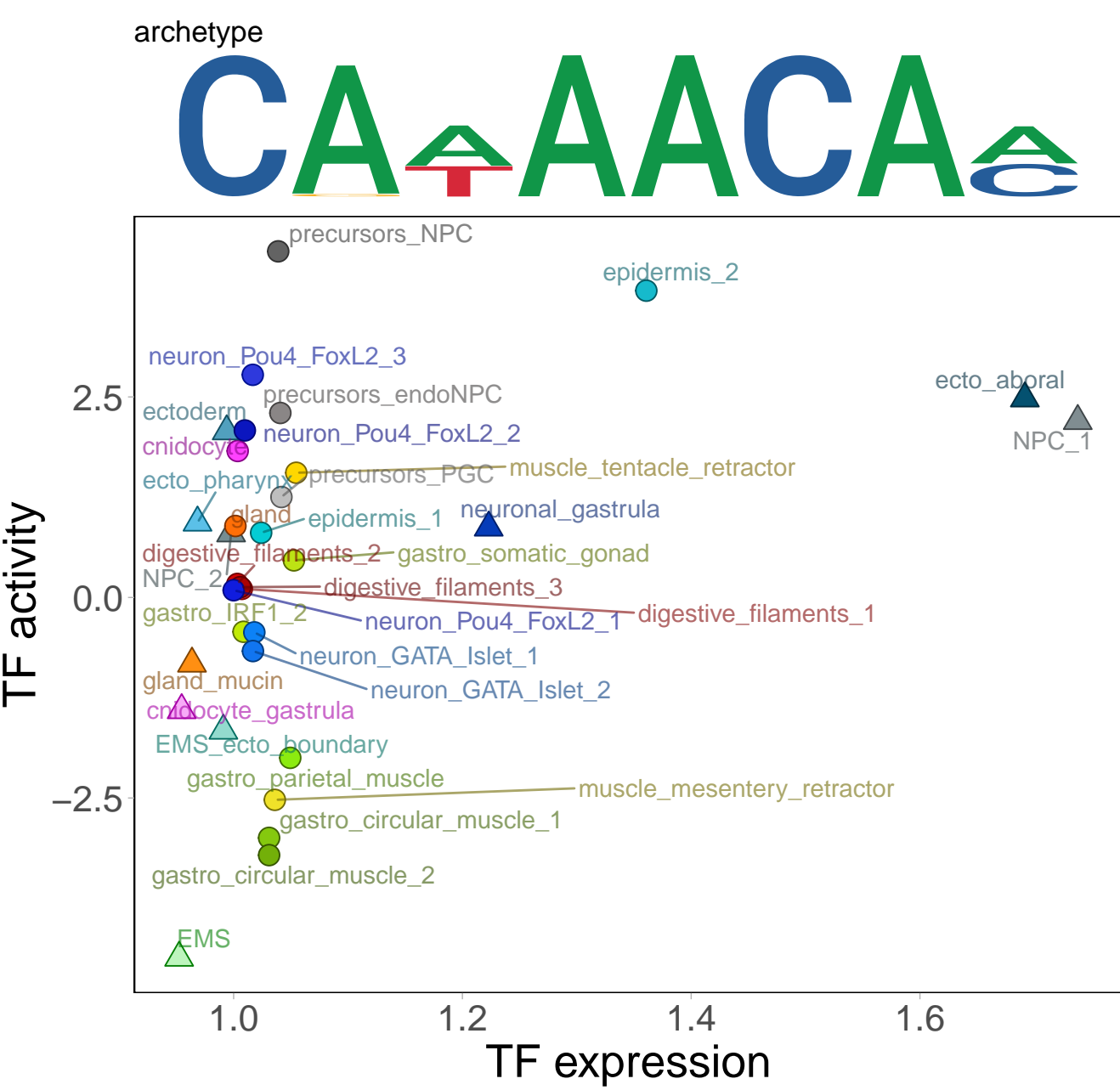
homer



streame

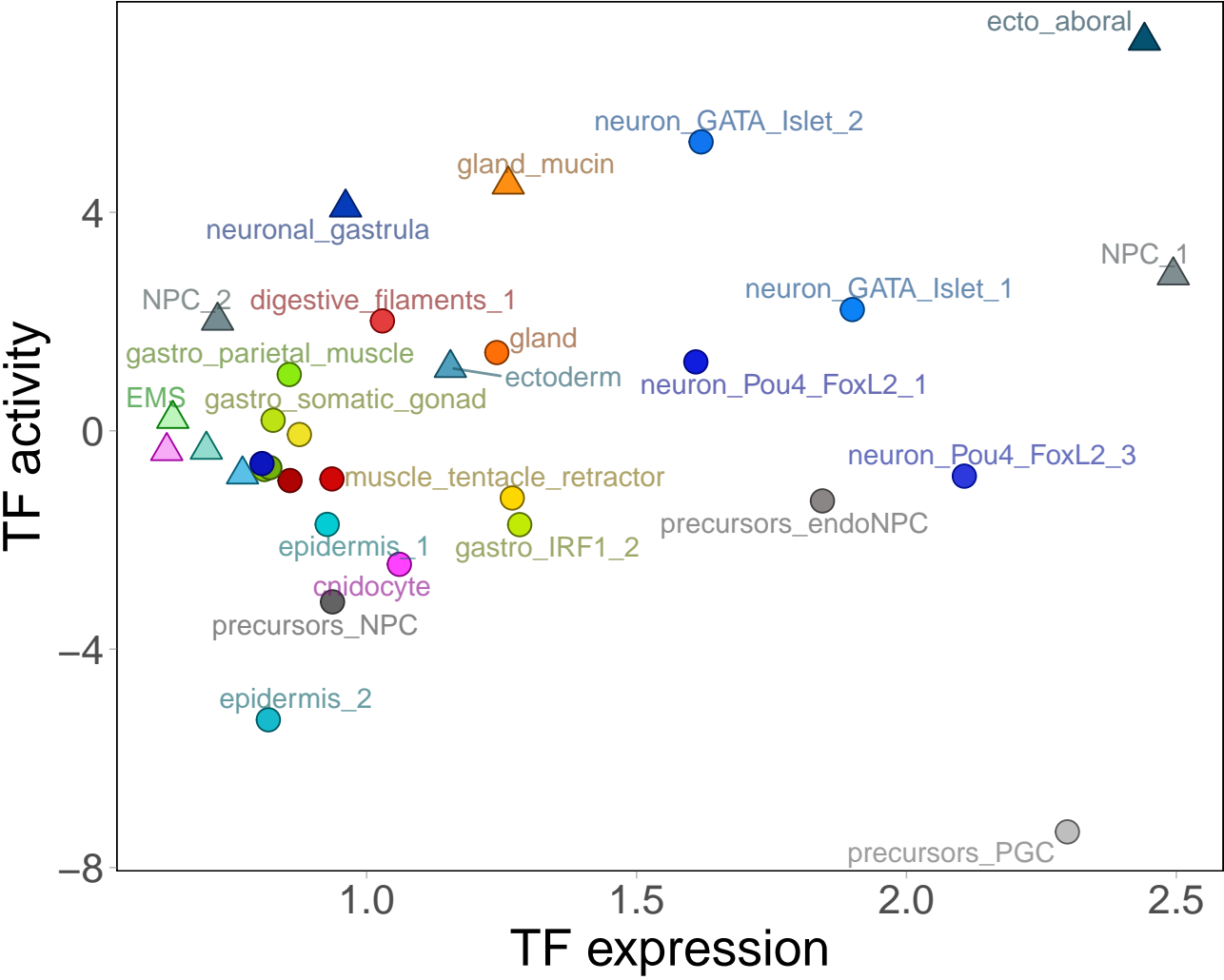


nvec_dap_17
Nvec_vc1.1_XM_001634111.3
OG: Forkhead.HG1.1:FOXA1/FOXA2/FOXA3/FOXB1/FOXB2/FOXD1/FOXD2/FOXD3/FOXD4/
PFAM: Forkhead

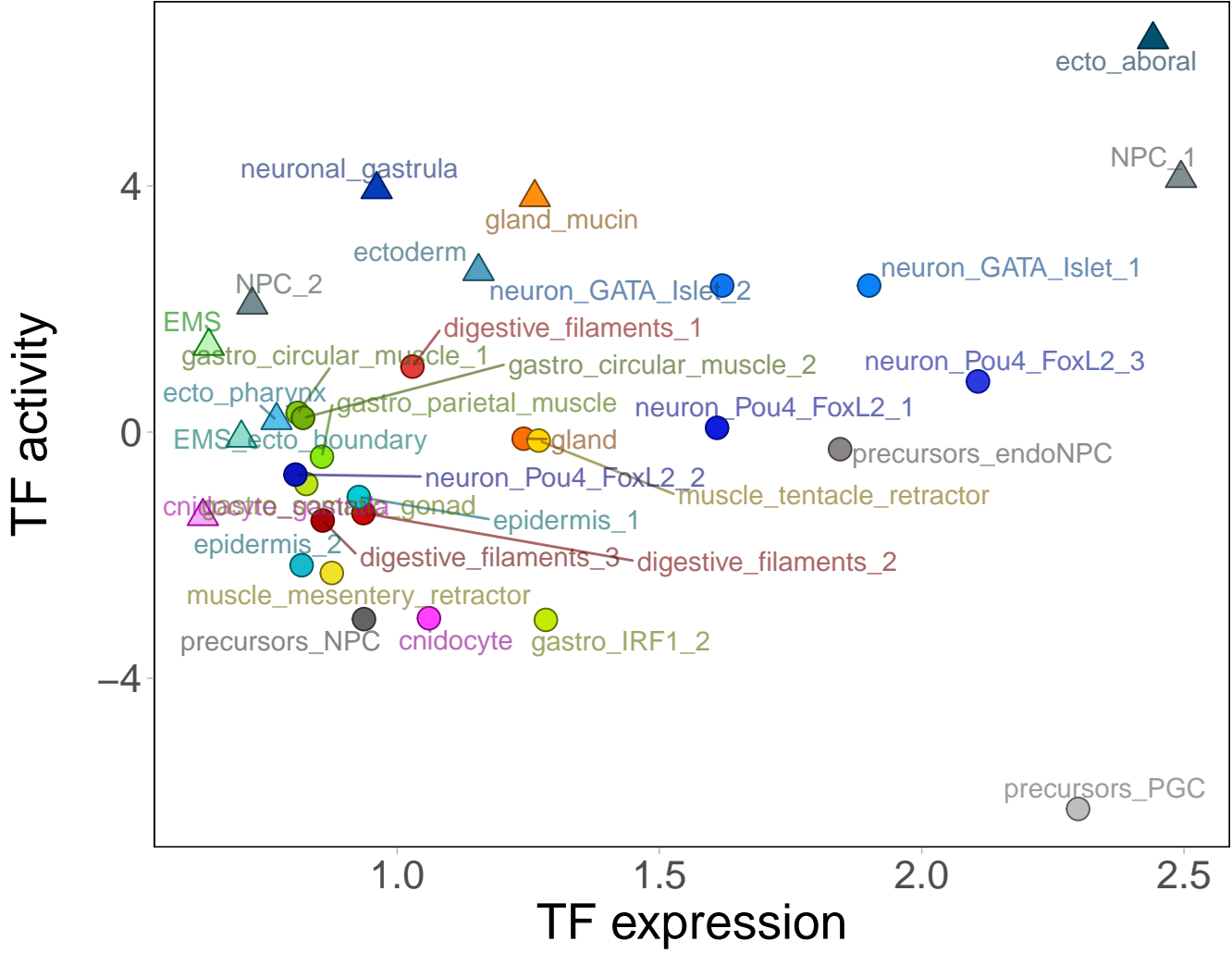


nvec_dap_21
Nvec_vc1.1_XM_032362713.2 (Islet)
OG: Homeodomains.HG2.19:ISL1/ISL2
PFAM: LIM/LIM/Homeodomain

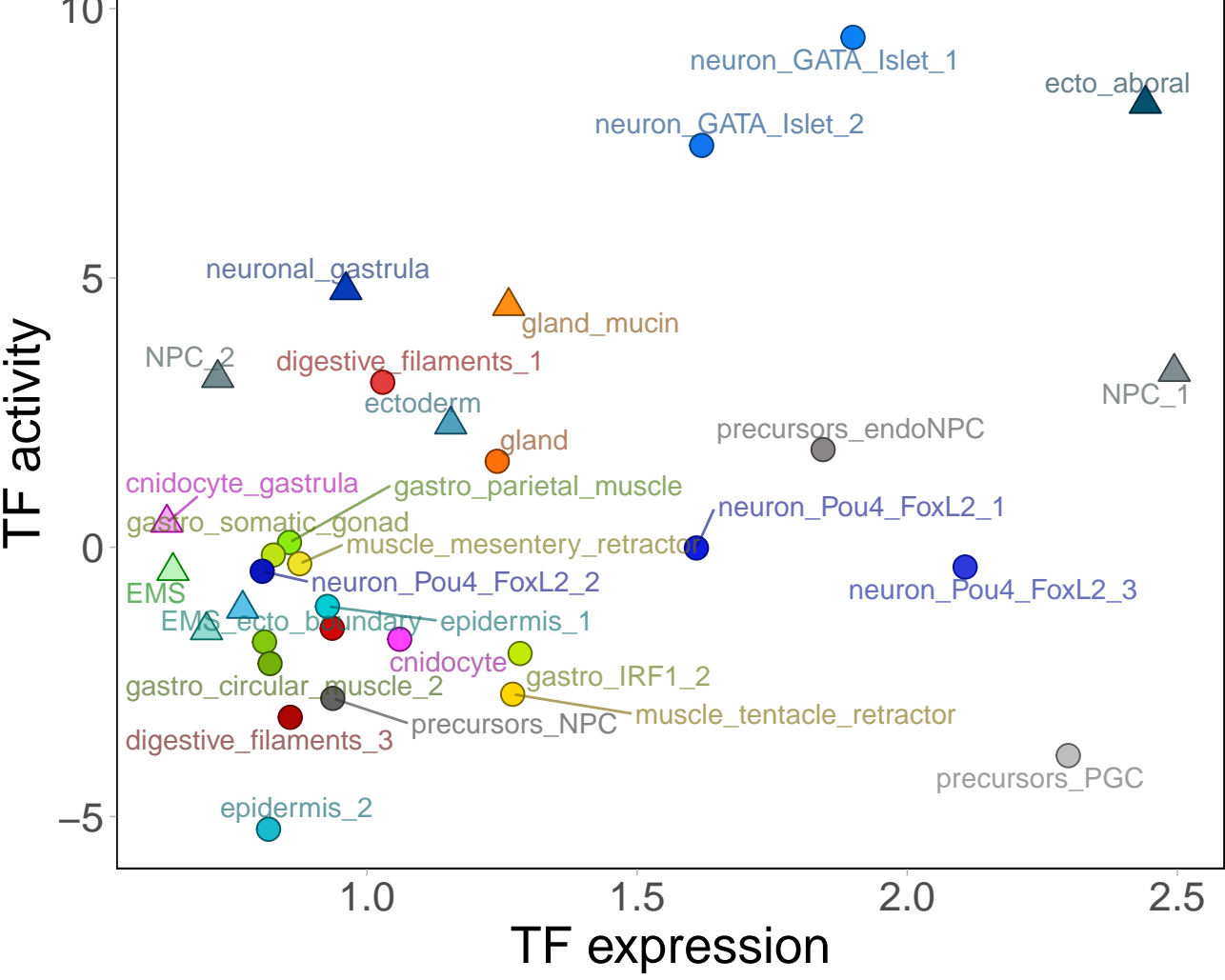
archetype



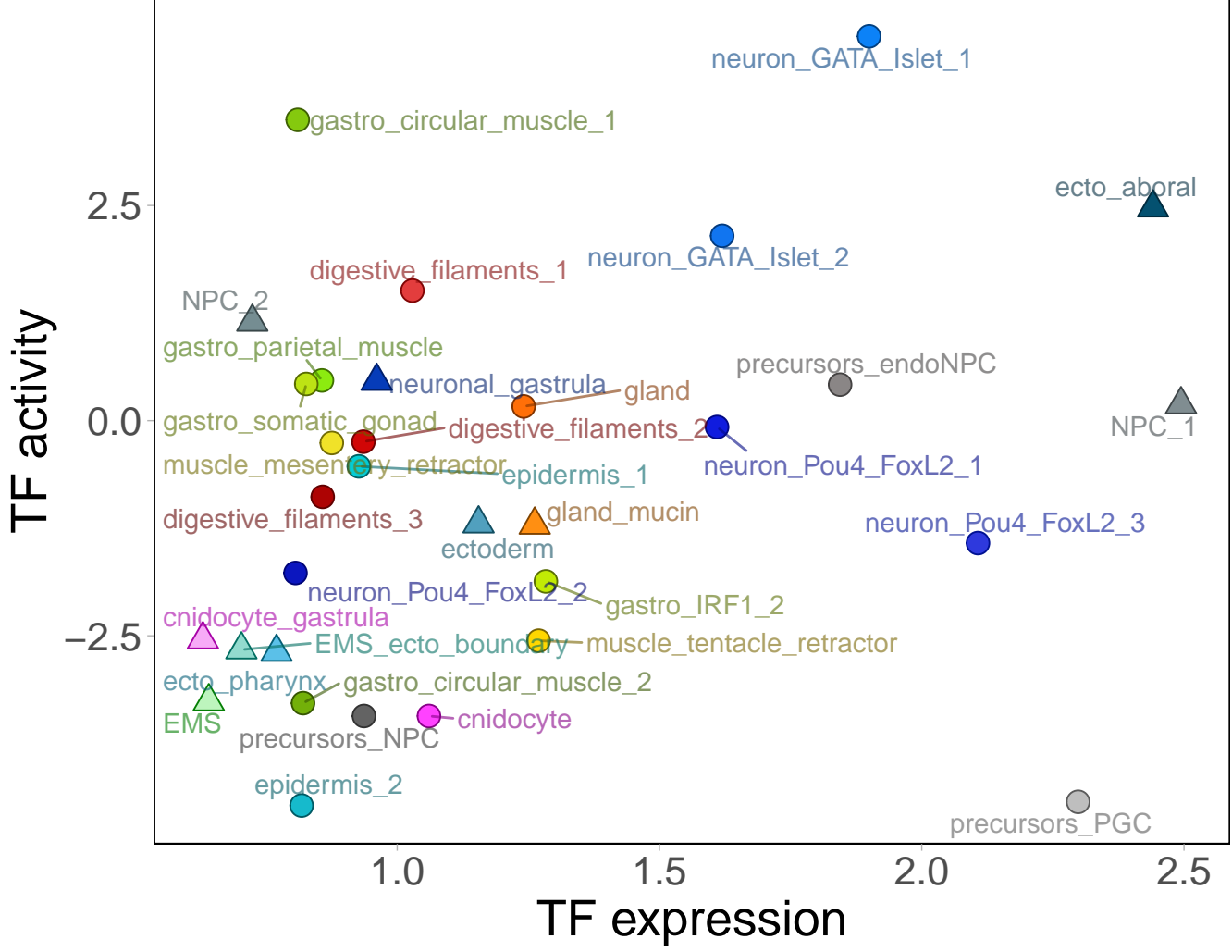
dimont



homer

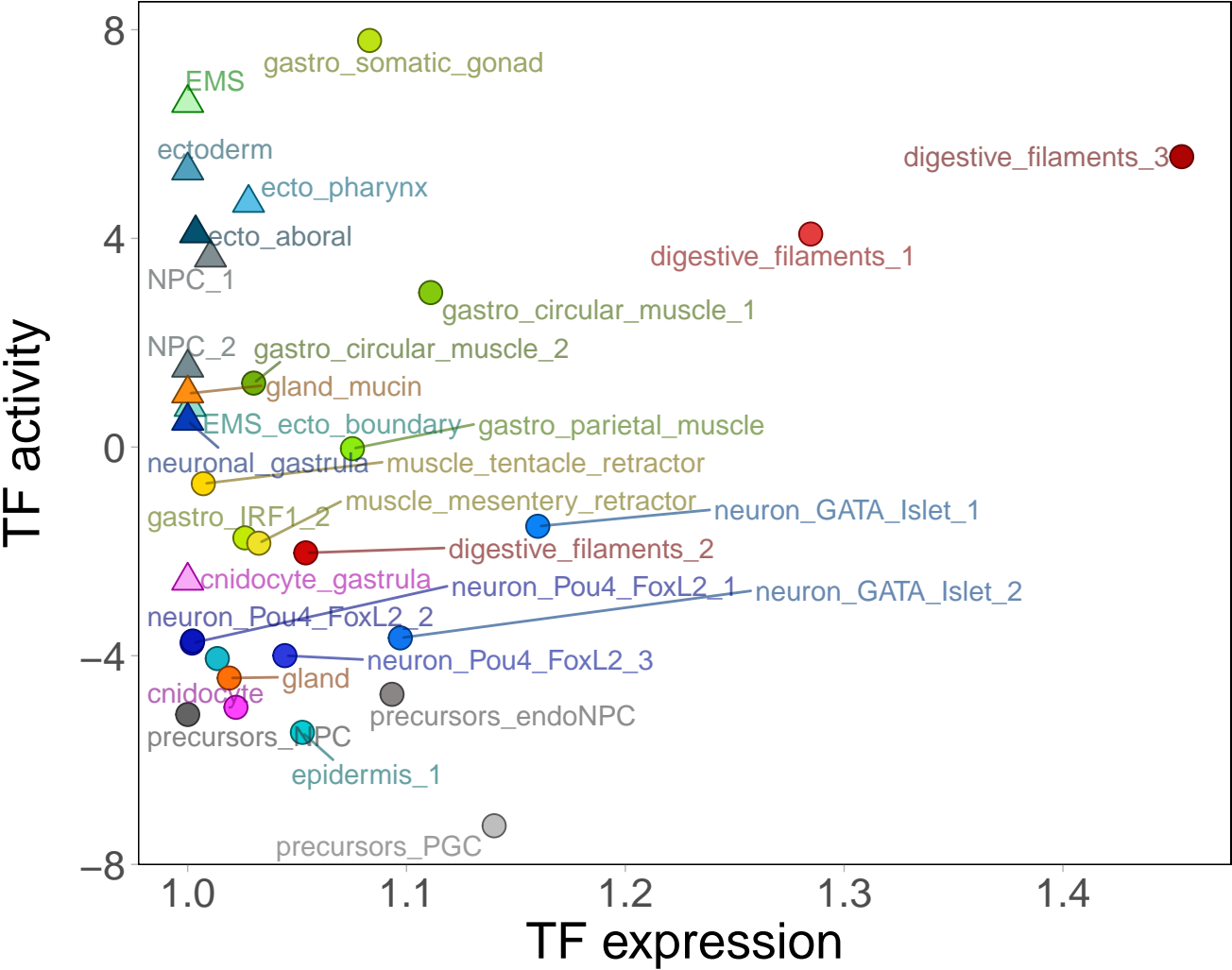


streame

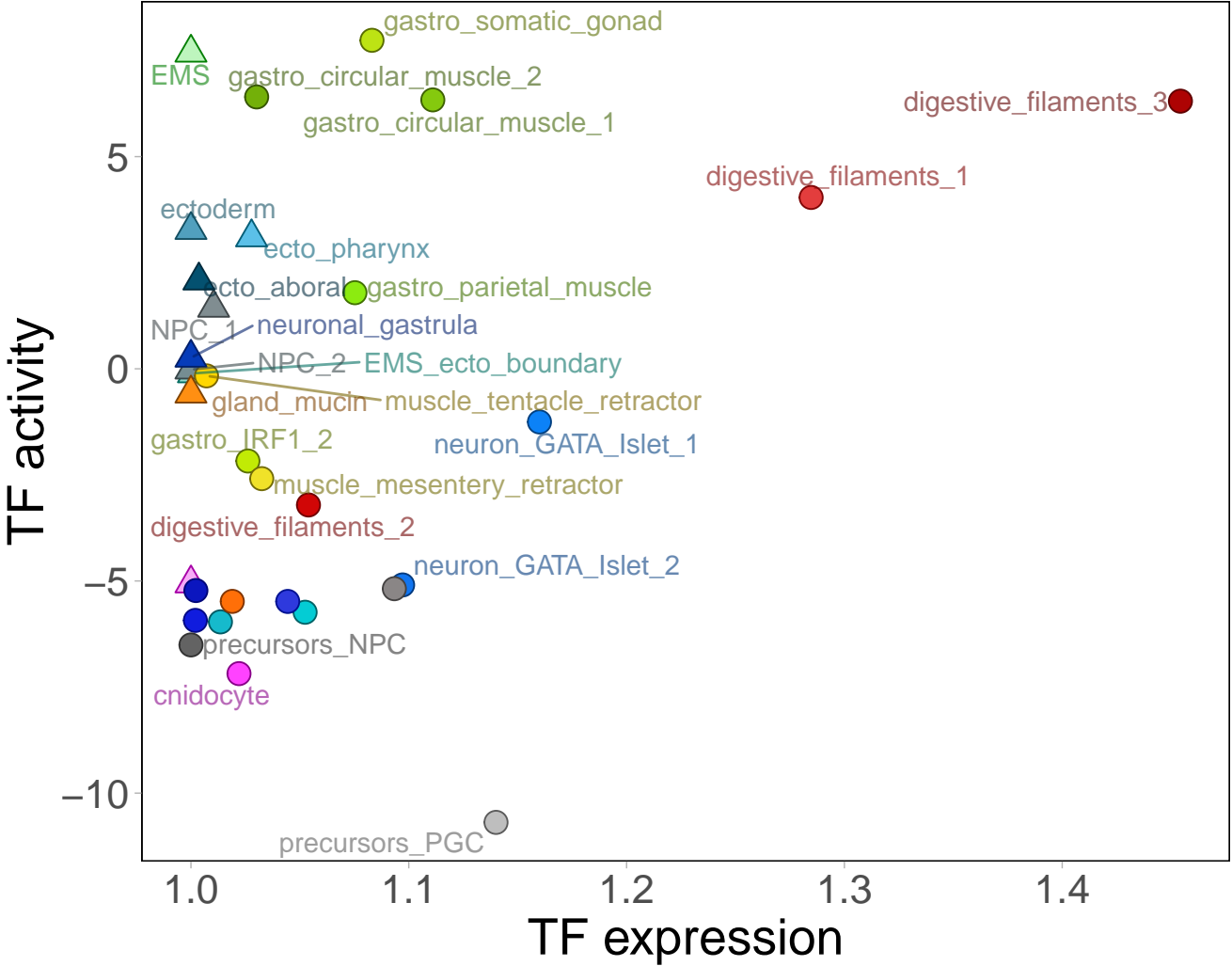


nvec_dap_22
Nvec_vc1.1_XM_032379400.2
OG: Homeodomains.HG2.76:like:DMBX1
PFAM: Homeodomain

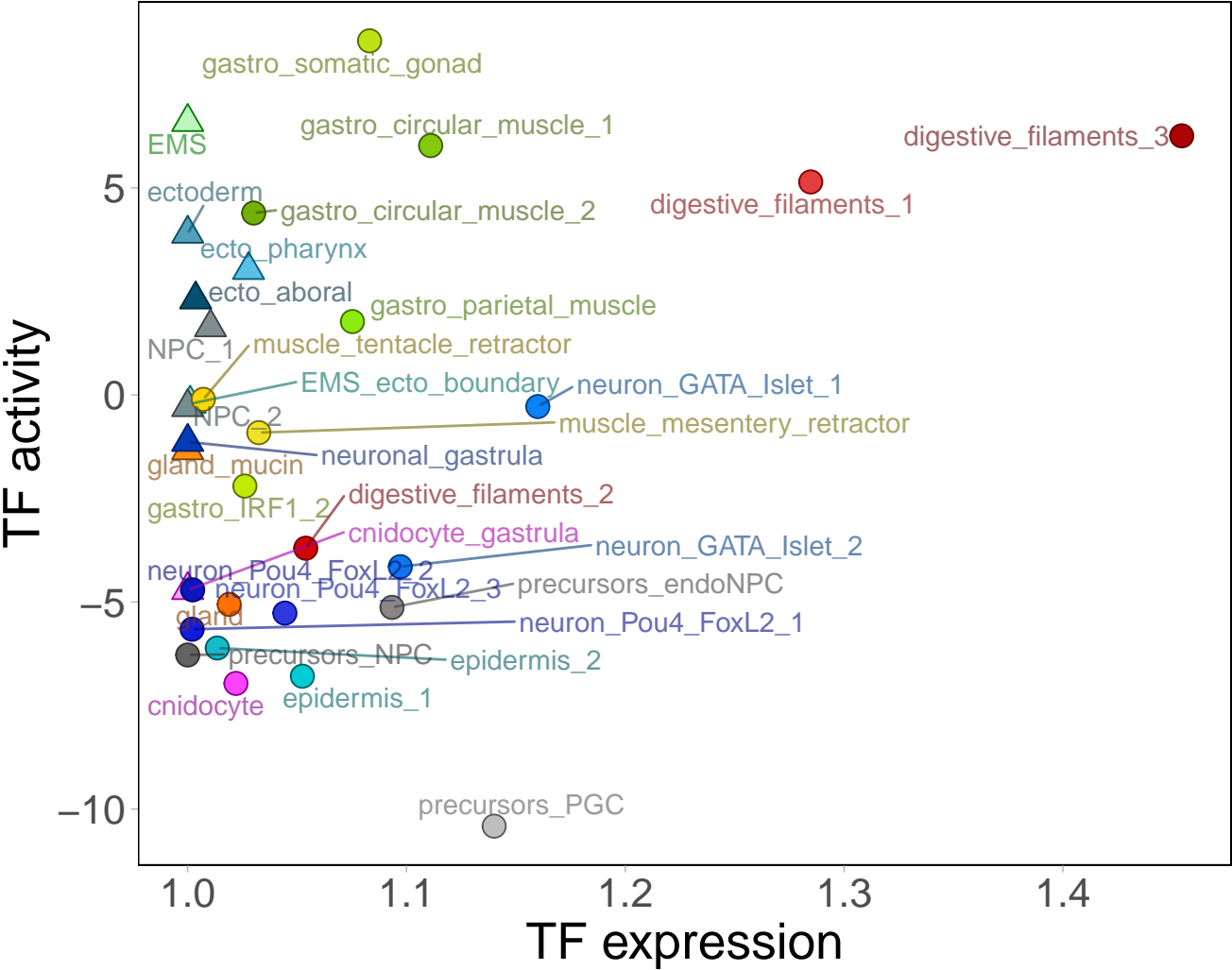
archetype



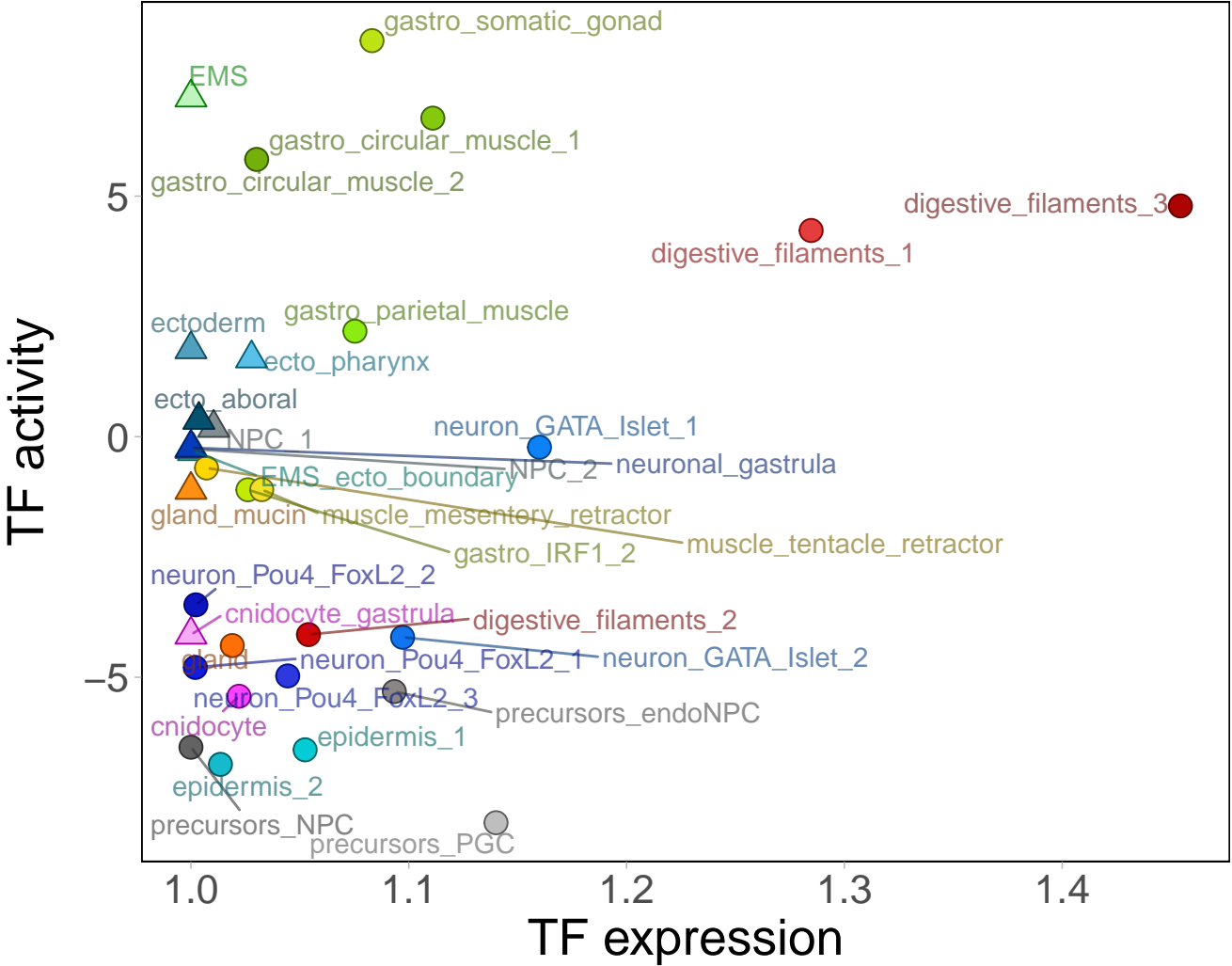
dimont



homer

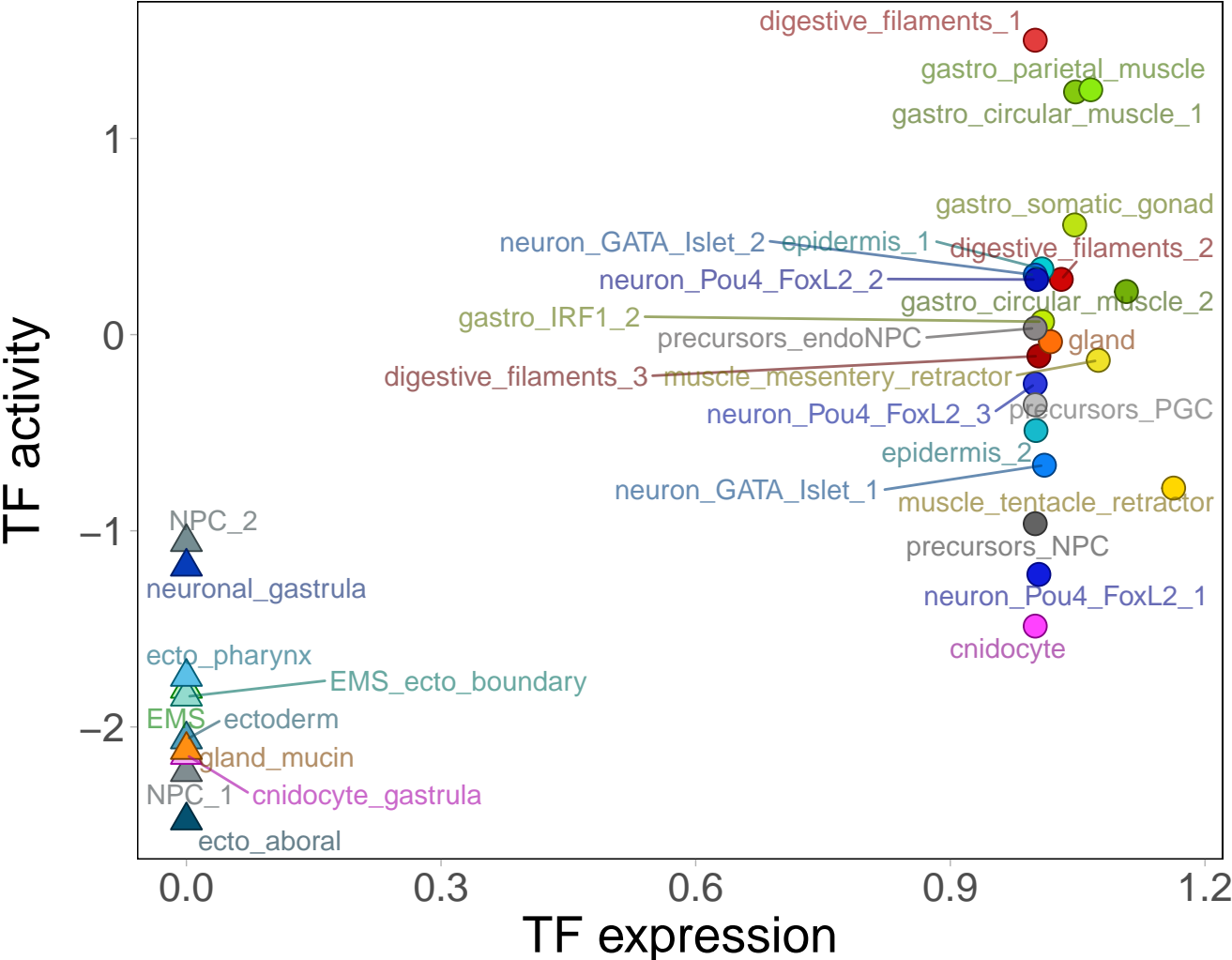


streme

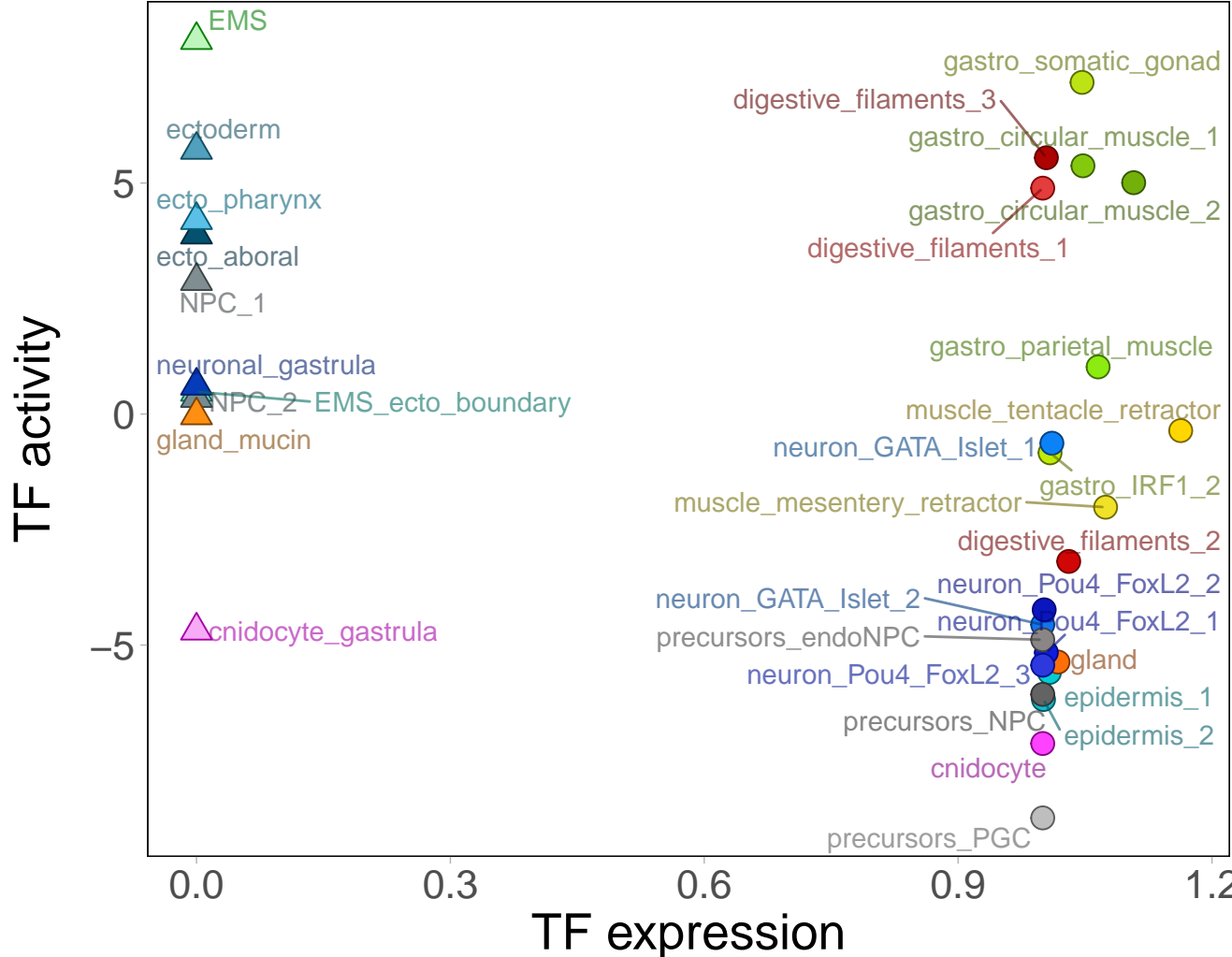


nvec_dap_24
Nvec_vc1.1_XM_001630676.3
OG: Homeodomains.HG2.68:like:ARX/OTP
PFAM: Homeodomain

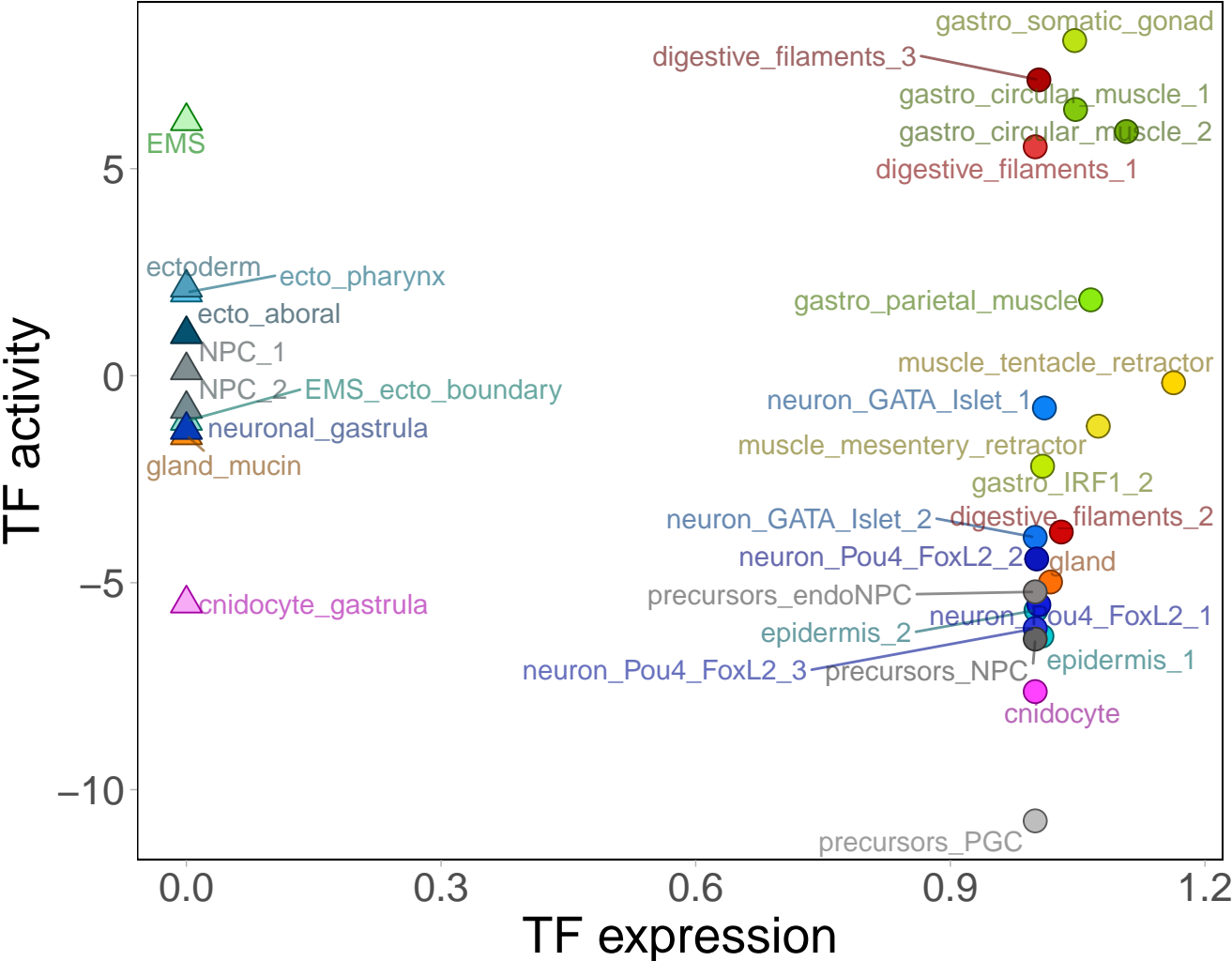
archetype



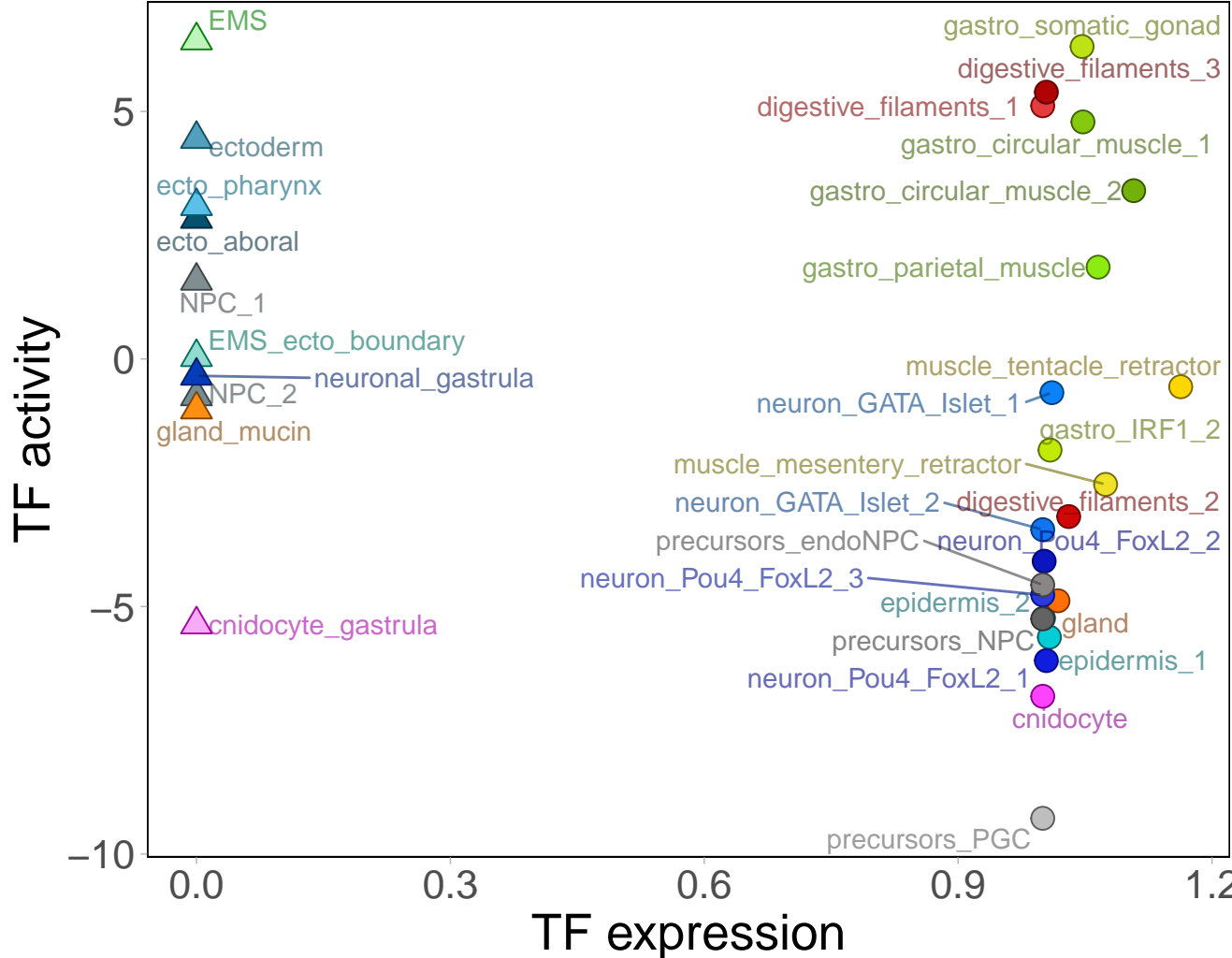
dimont



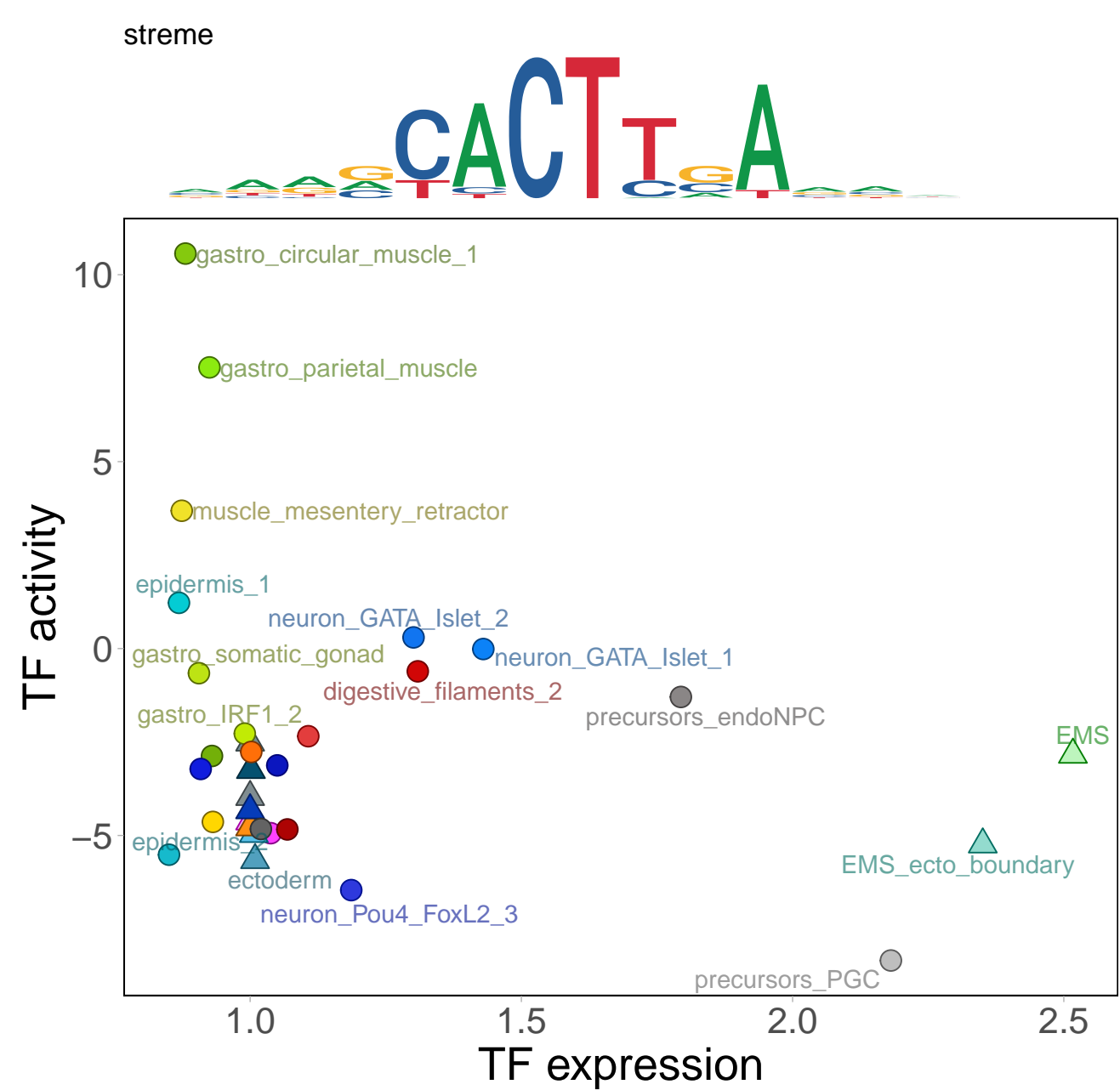
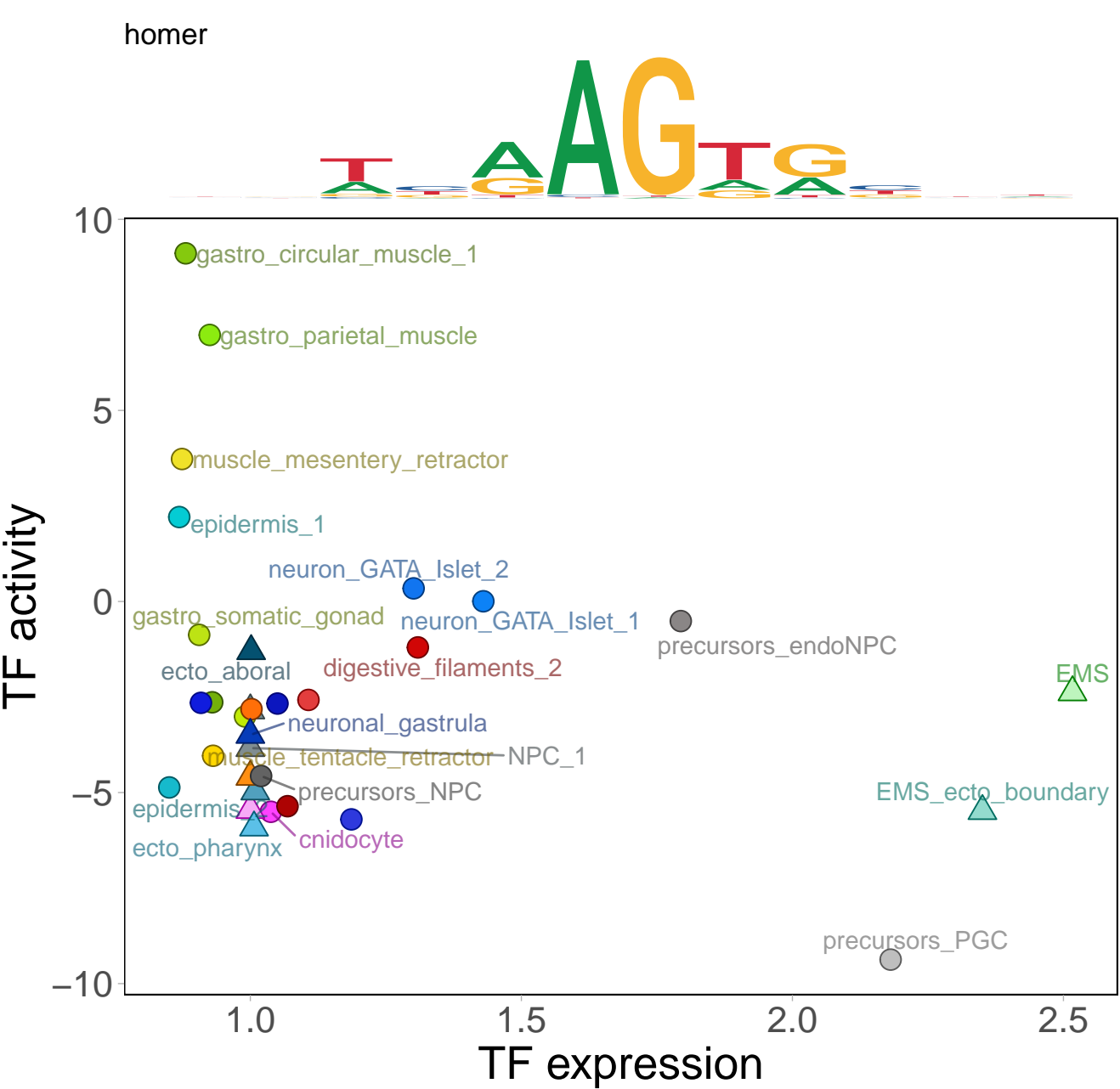
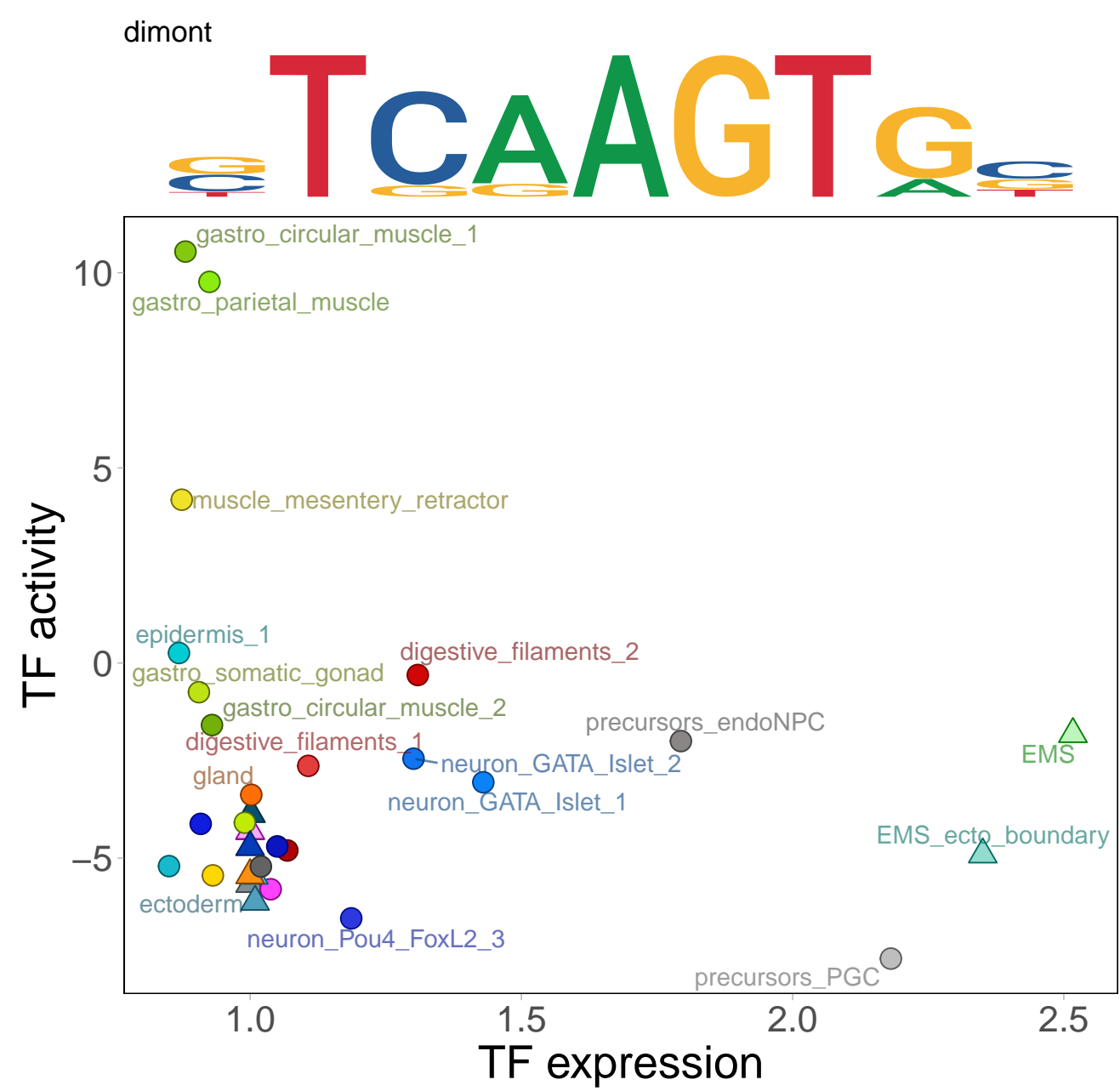
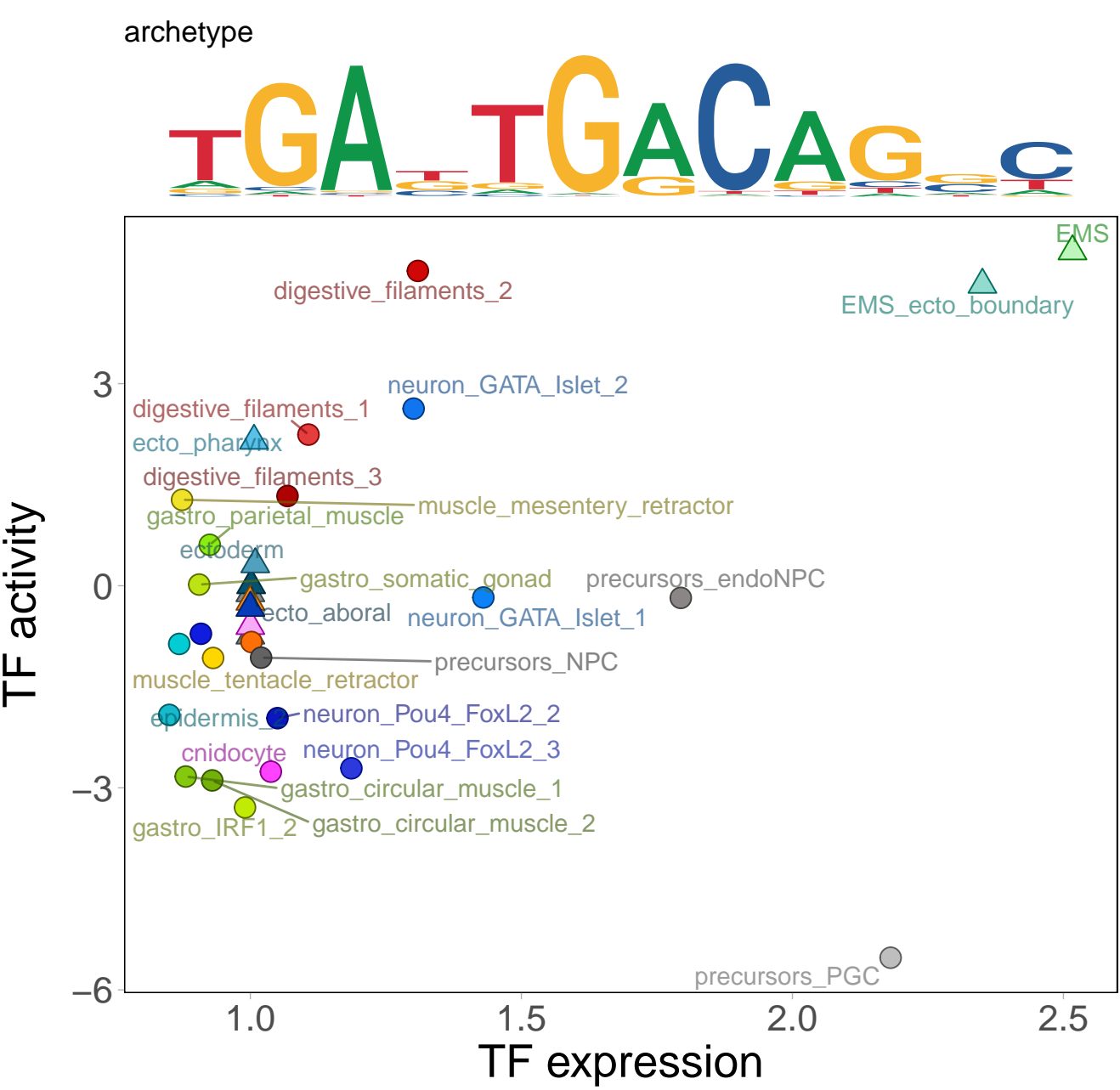
homer



streme

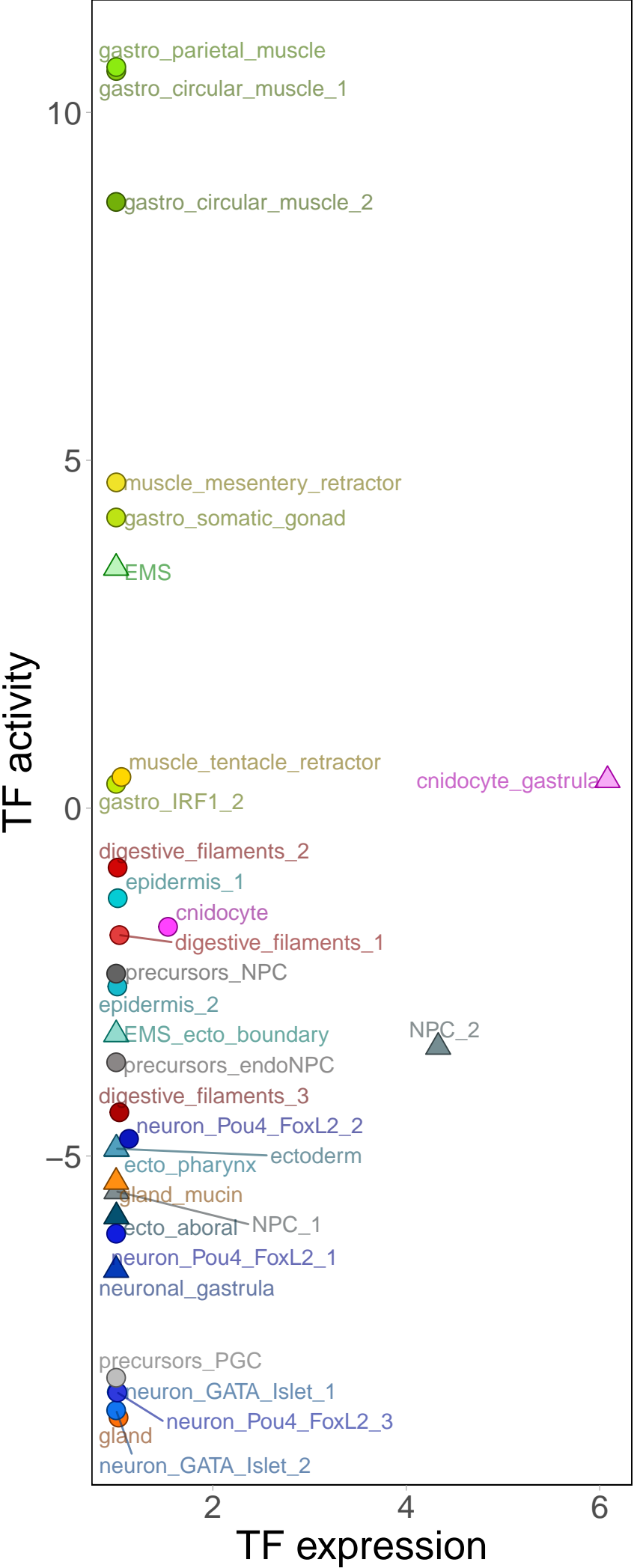


nvec_dap_25
Nvec_vc1.1_XM_032373618.2 (Nkx-2.2a)
OG: Homeodomains.HG1.19:NKX2-1/NKX2-2/NKX2-3/NKX2-4/NKX2-5/NKX2-6/NKX2-8/M
PFAM: Homeodomain

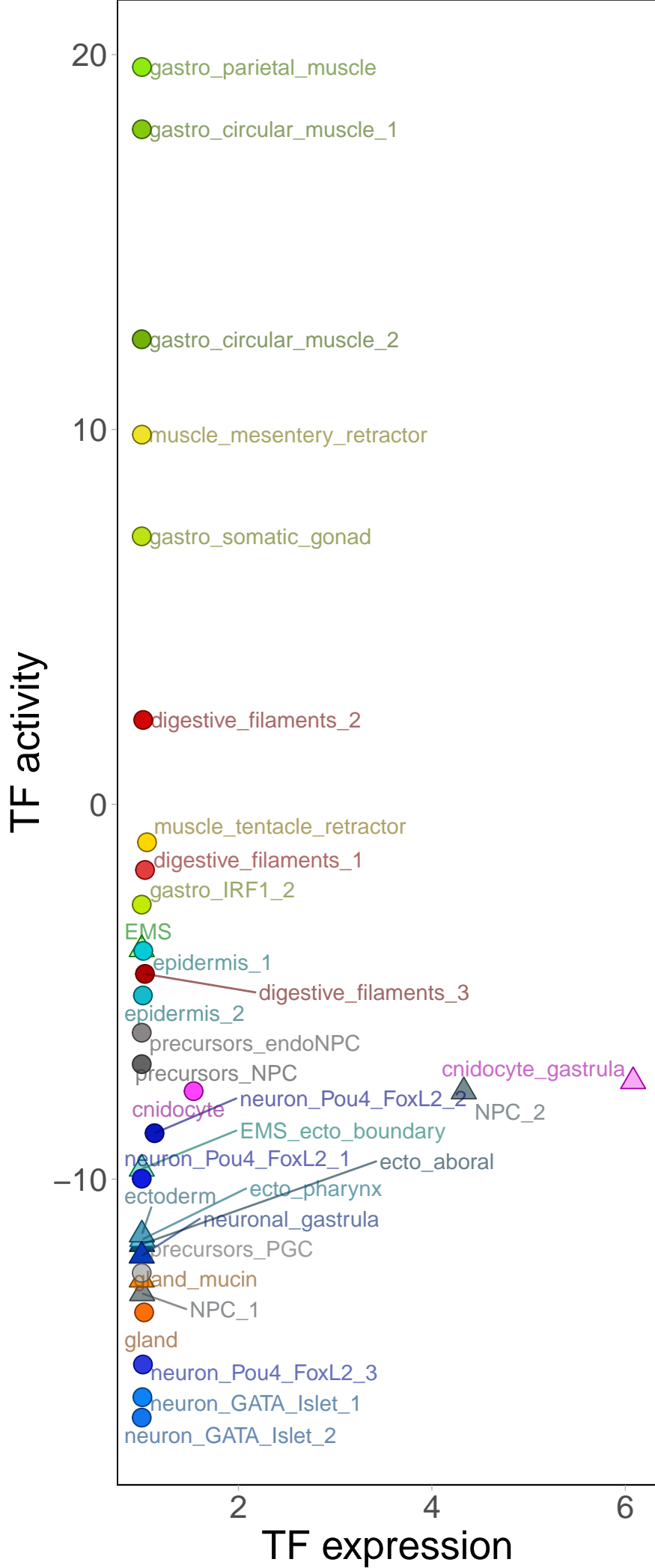
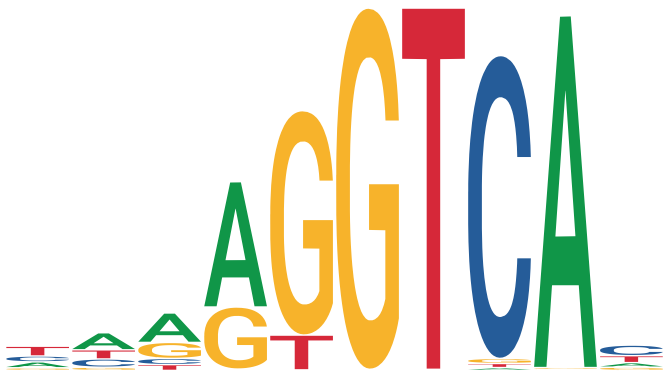


nvec_dap_27
Nvec_vc1.1_XM_001634328.3 (NR12)
OG: zf-C4_Nuclear_receptors.HG1.24:like:NR2F1/NR2F2/NR2F6
PFAM: zf-C4/Hormone_recep

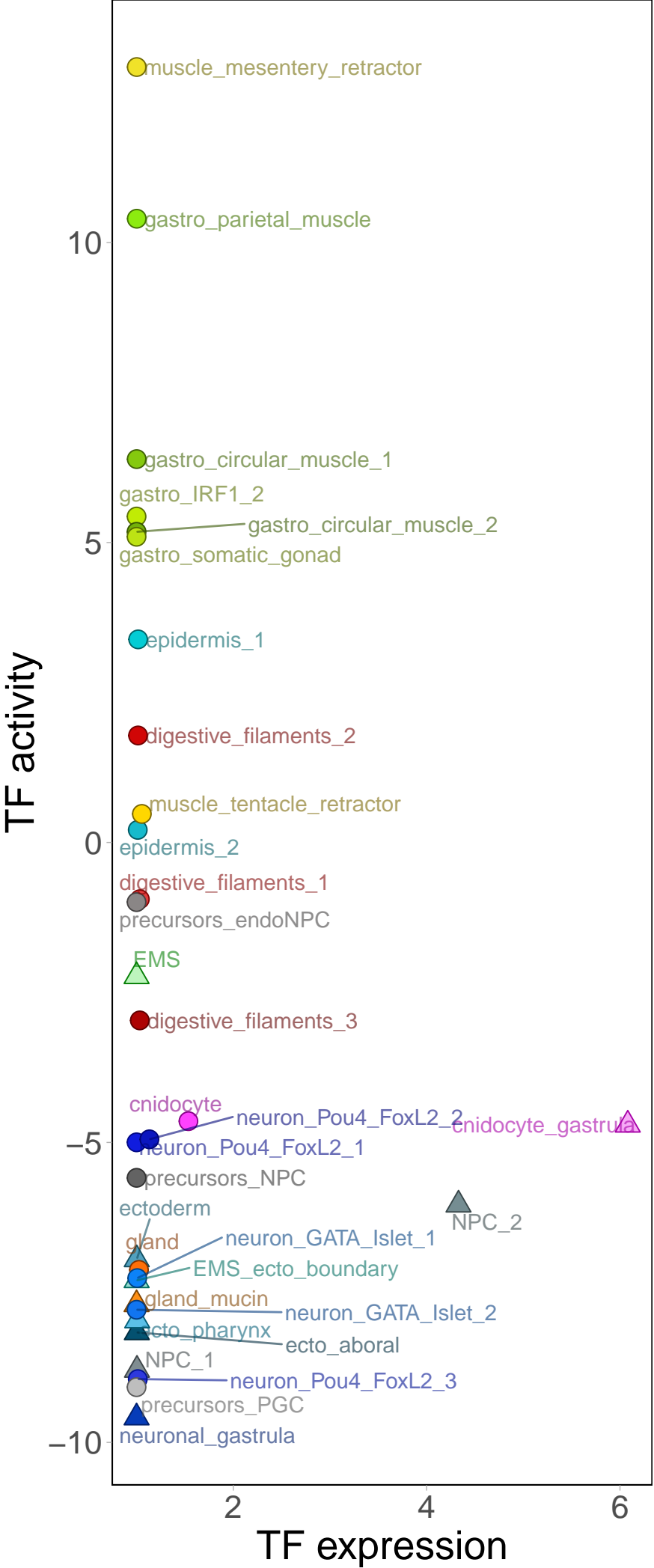
dimont



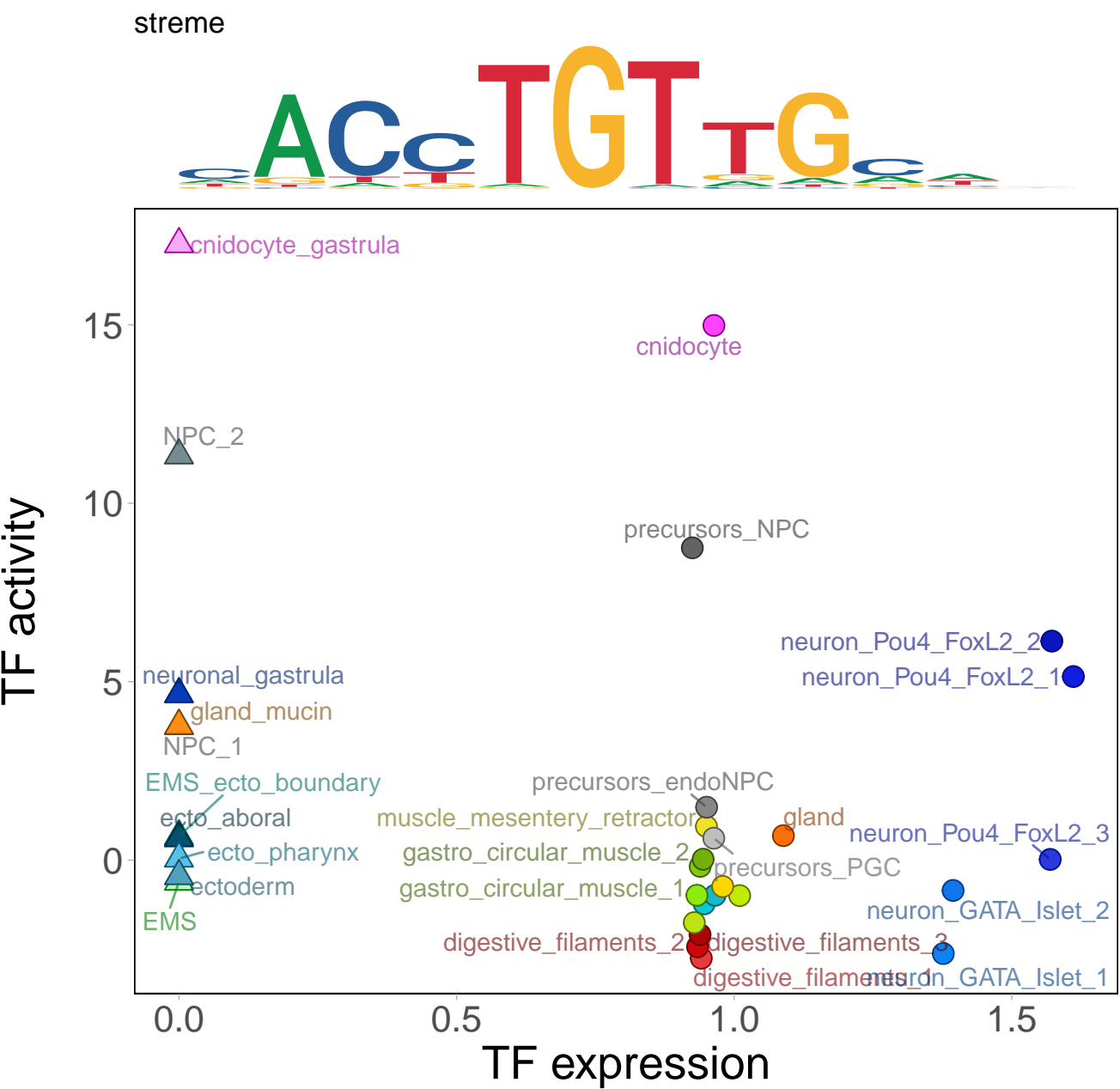
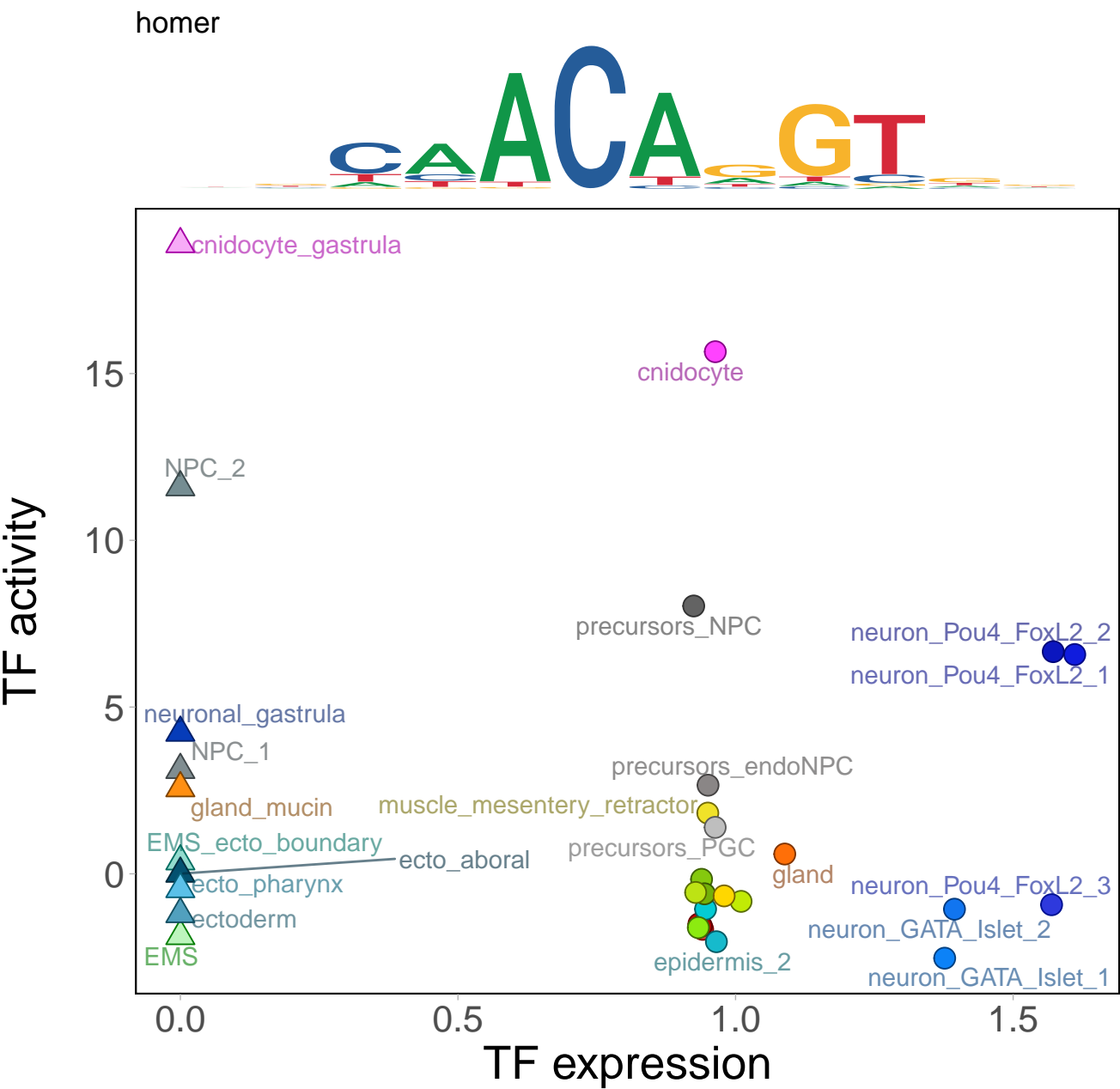
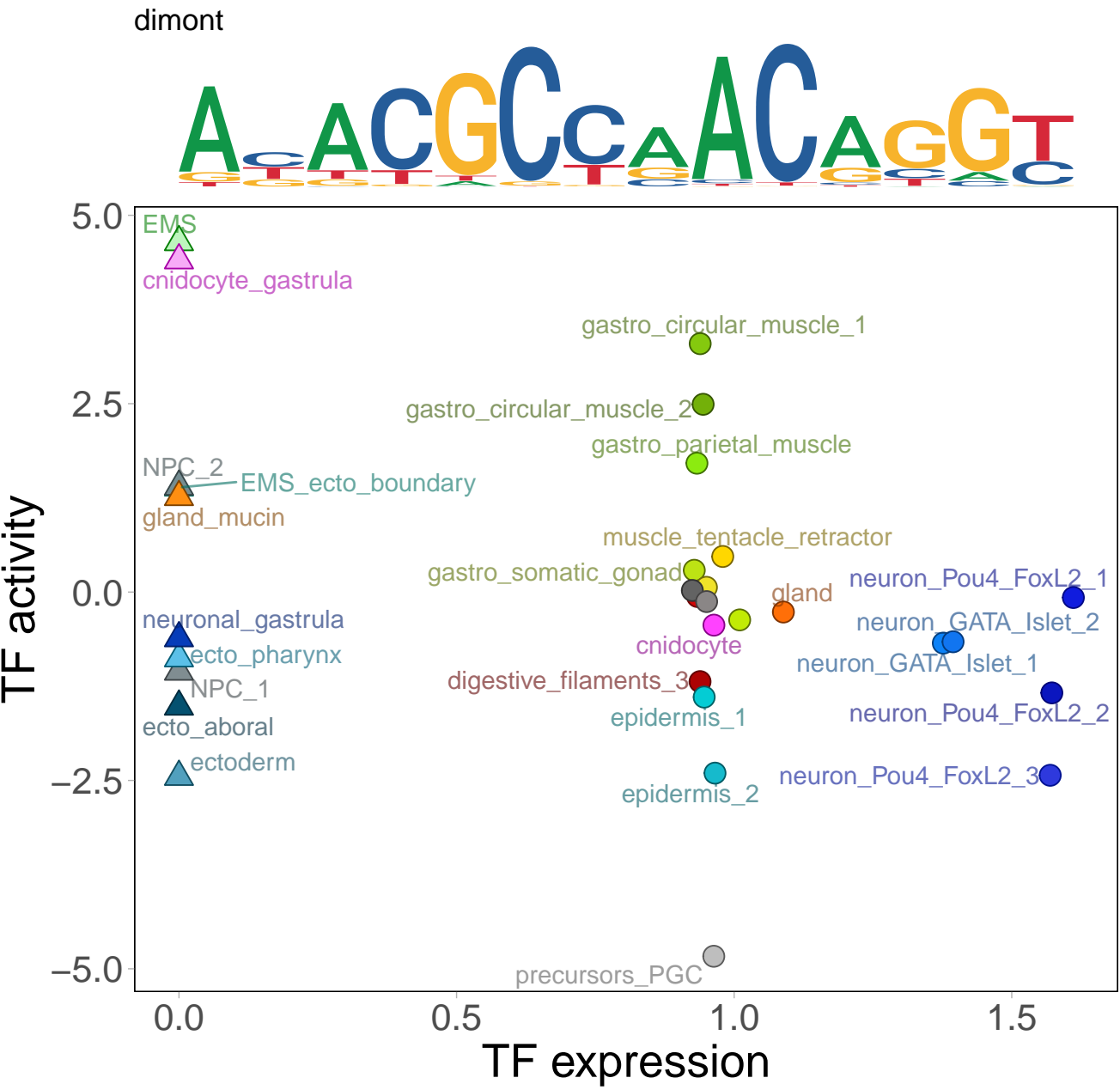
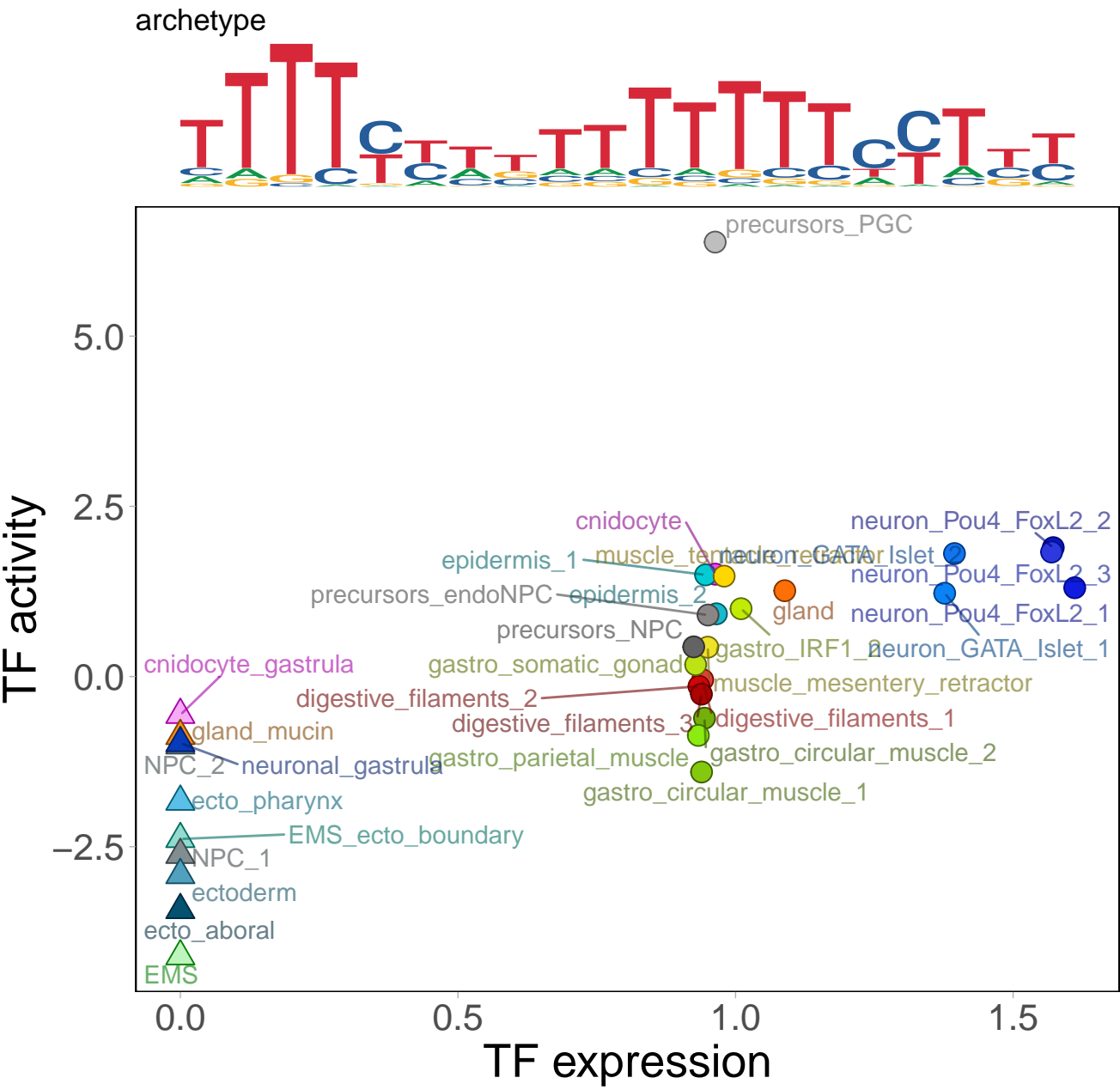
homer



streme

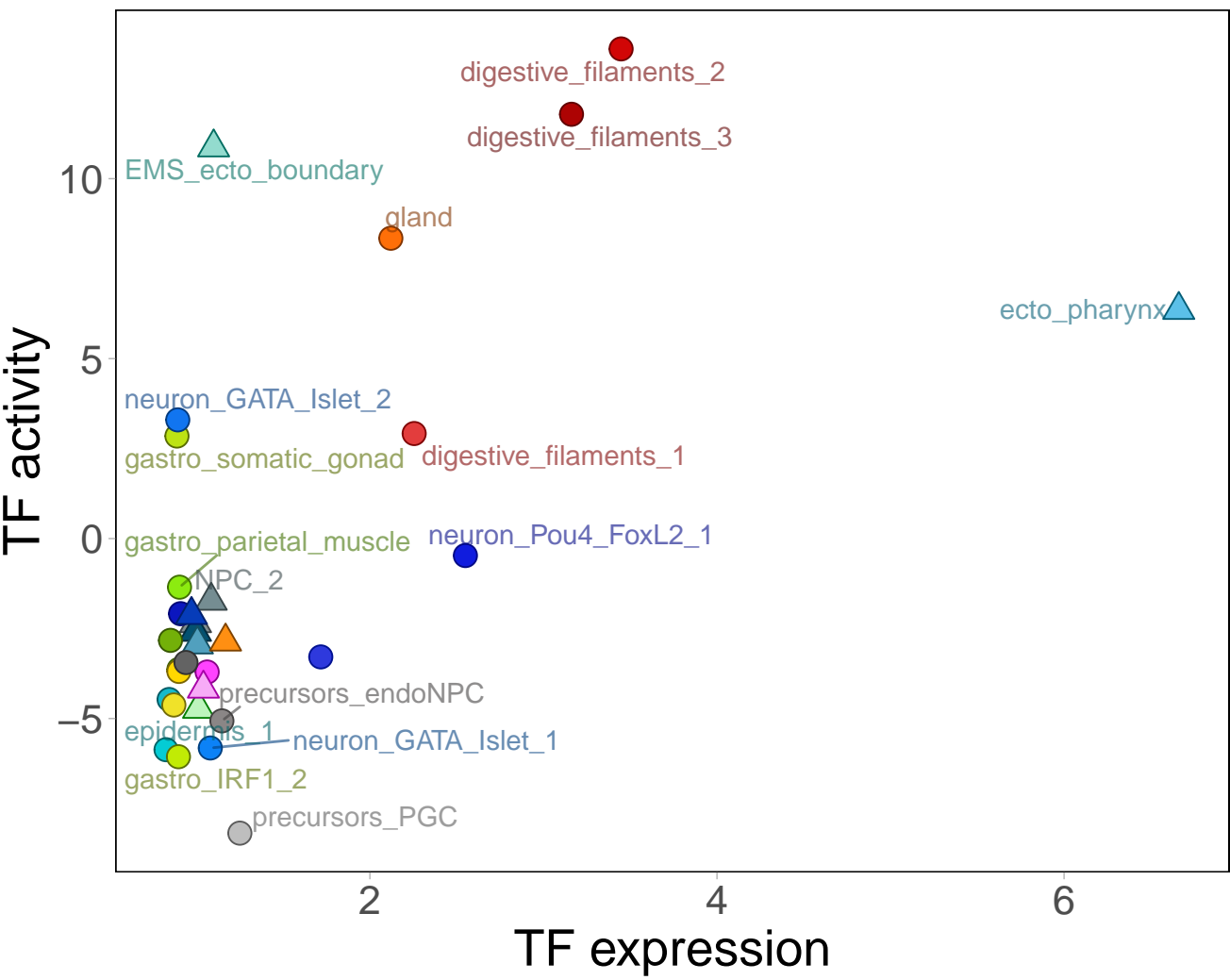


nvec_dap_29
Nvec_vc1.1_XM_032381270.2
OG: zf-C2H2.HG7.4:SCRT1/SCRT2
PFAM: zf-C2H2/zf-C2H2/zf-C2H2/zf-C2H2

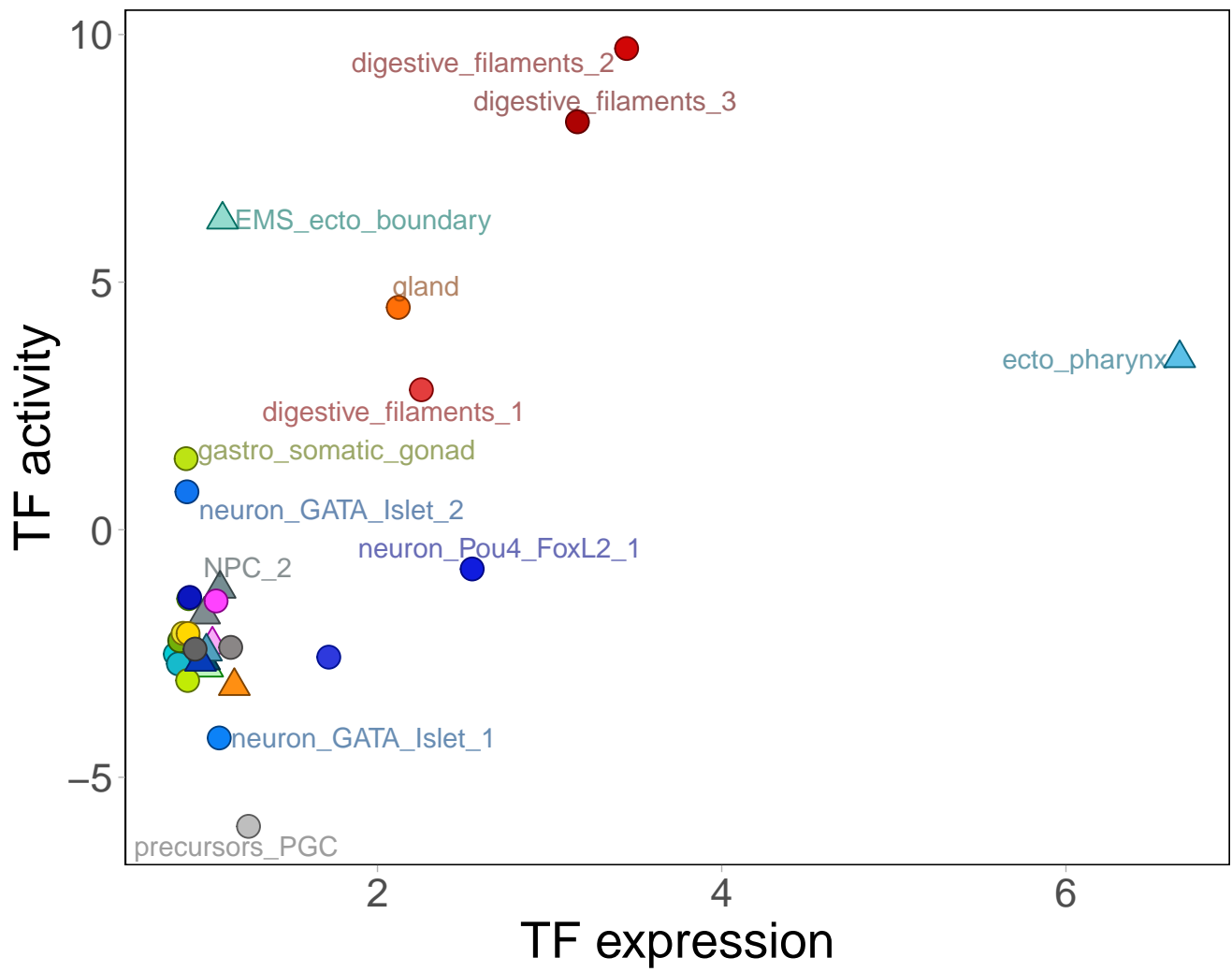


nvec_dap_30
Nvec_vc1.1_XM_001634505.3 (FoxA)
OG: Forkhead.HG1.1:FOXA1/FOXA2/FOXA3/FOXB1/FOXB2/FOXD1/FOXD2/FOXD3/FOXD4/
PFAM: Forkhead_N/Forkhead/HNF_C

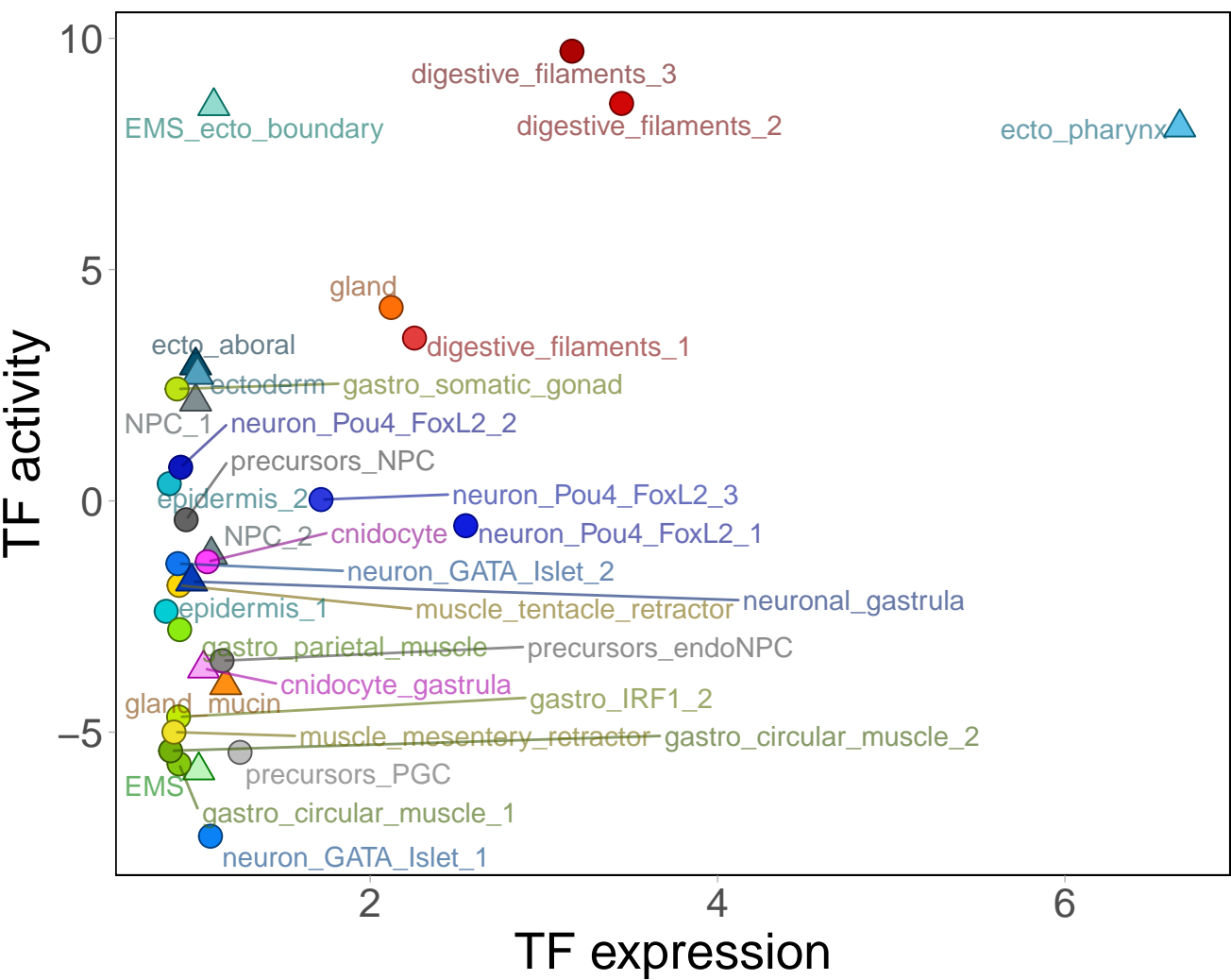
archetype



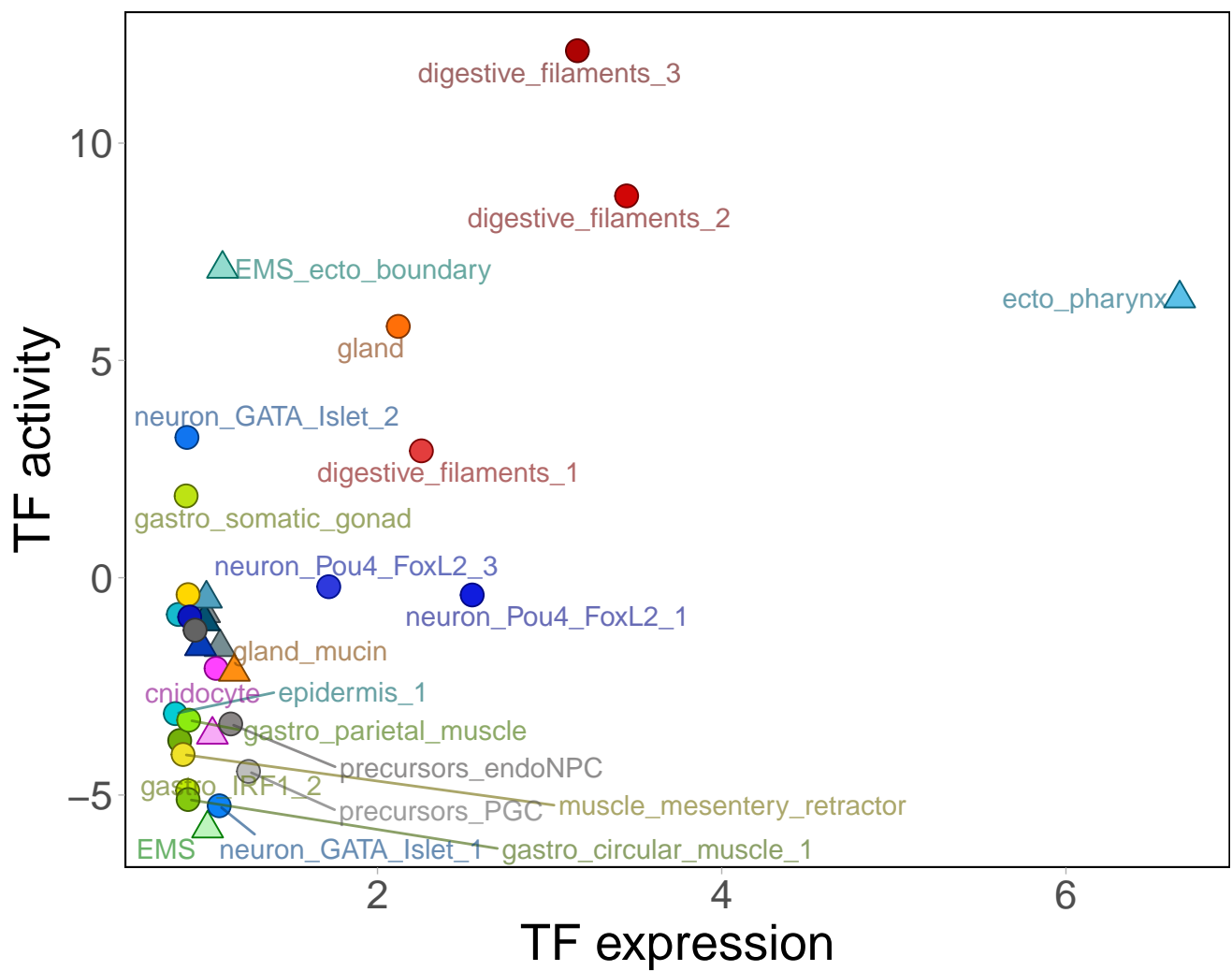
dimont



homer

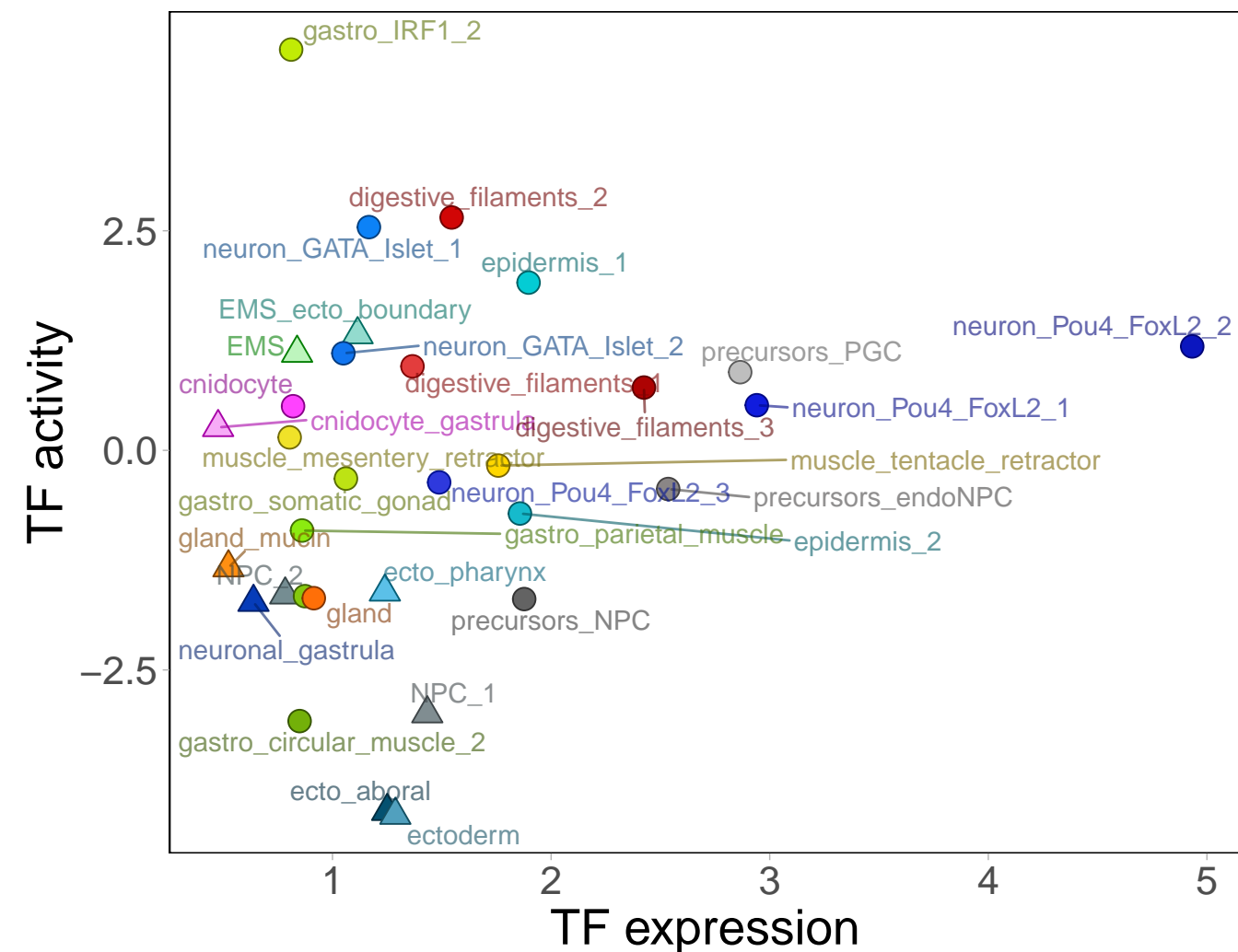


streame

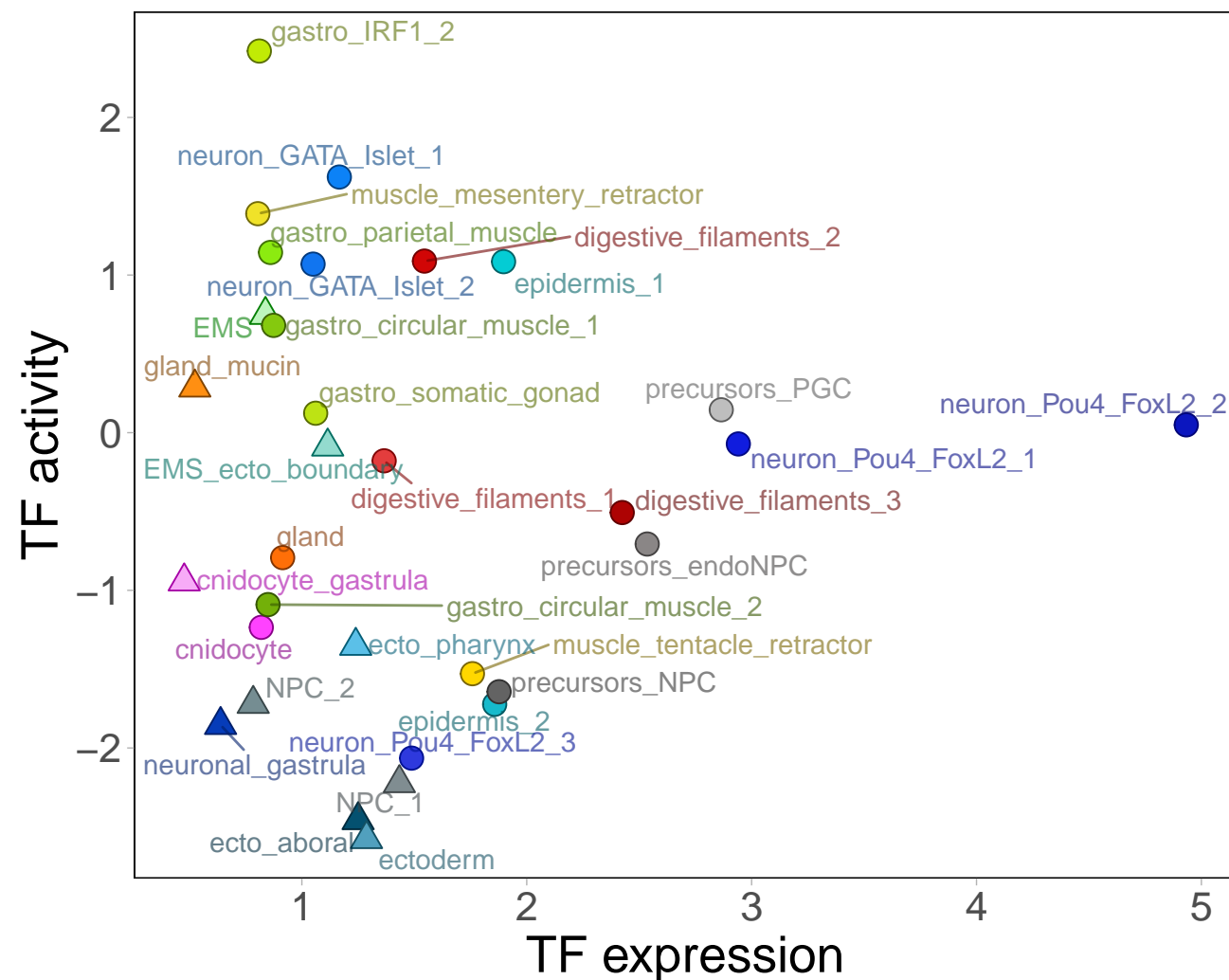
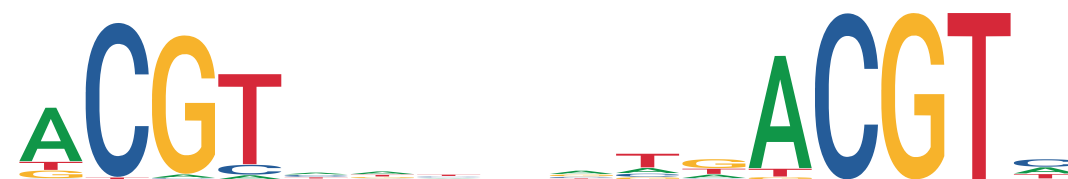


nvec_dap_33
Nvec_vc1.1_XM_001623467.3
OG: bZIP.HG1.12:CREBZF
PFAM: bZIP_2

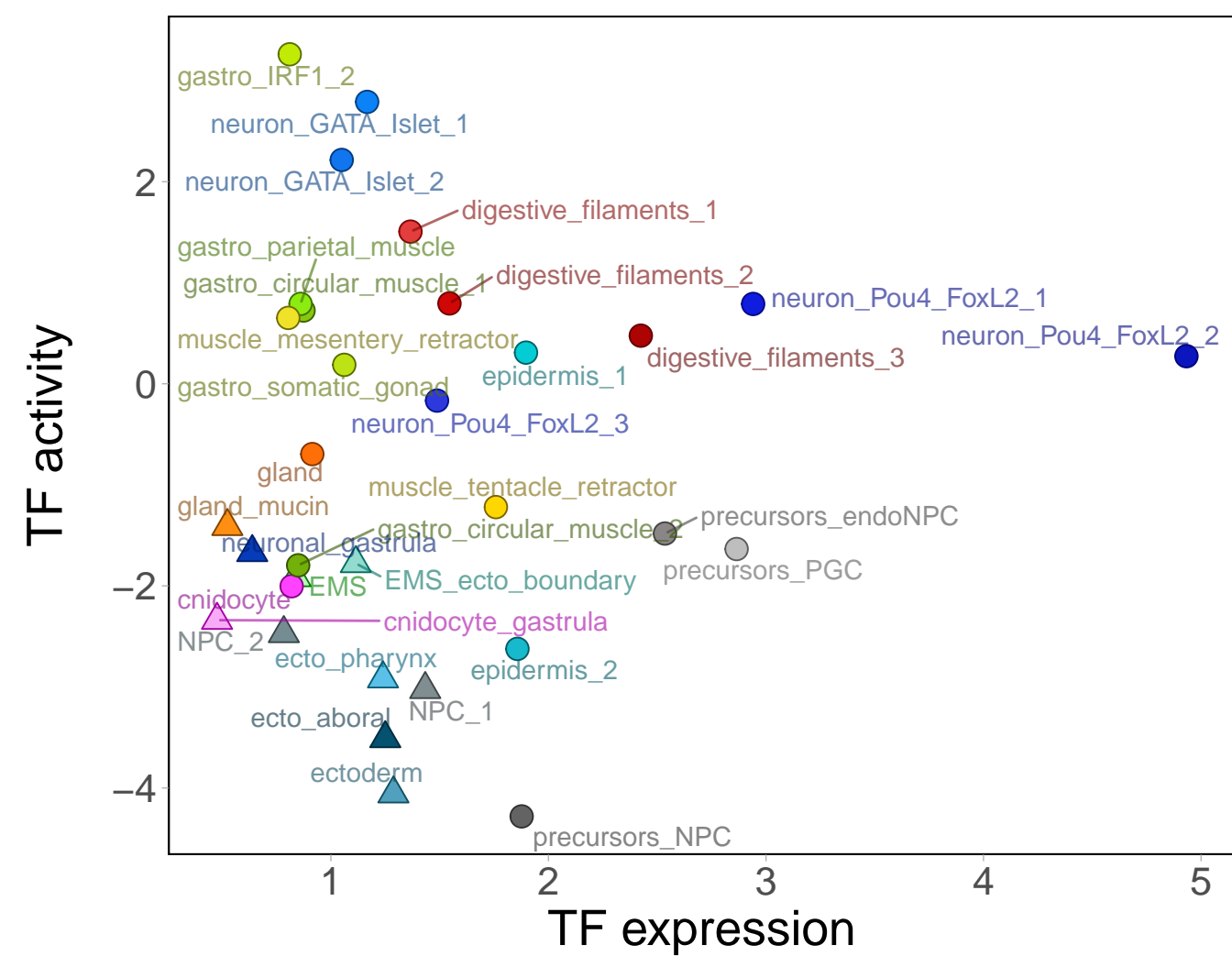
archetype



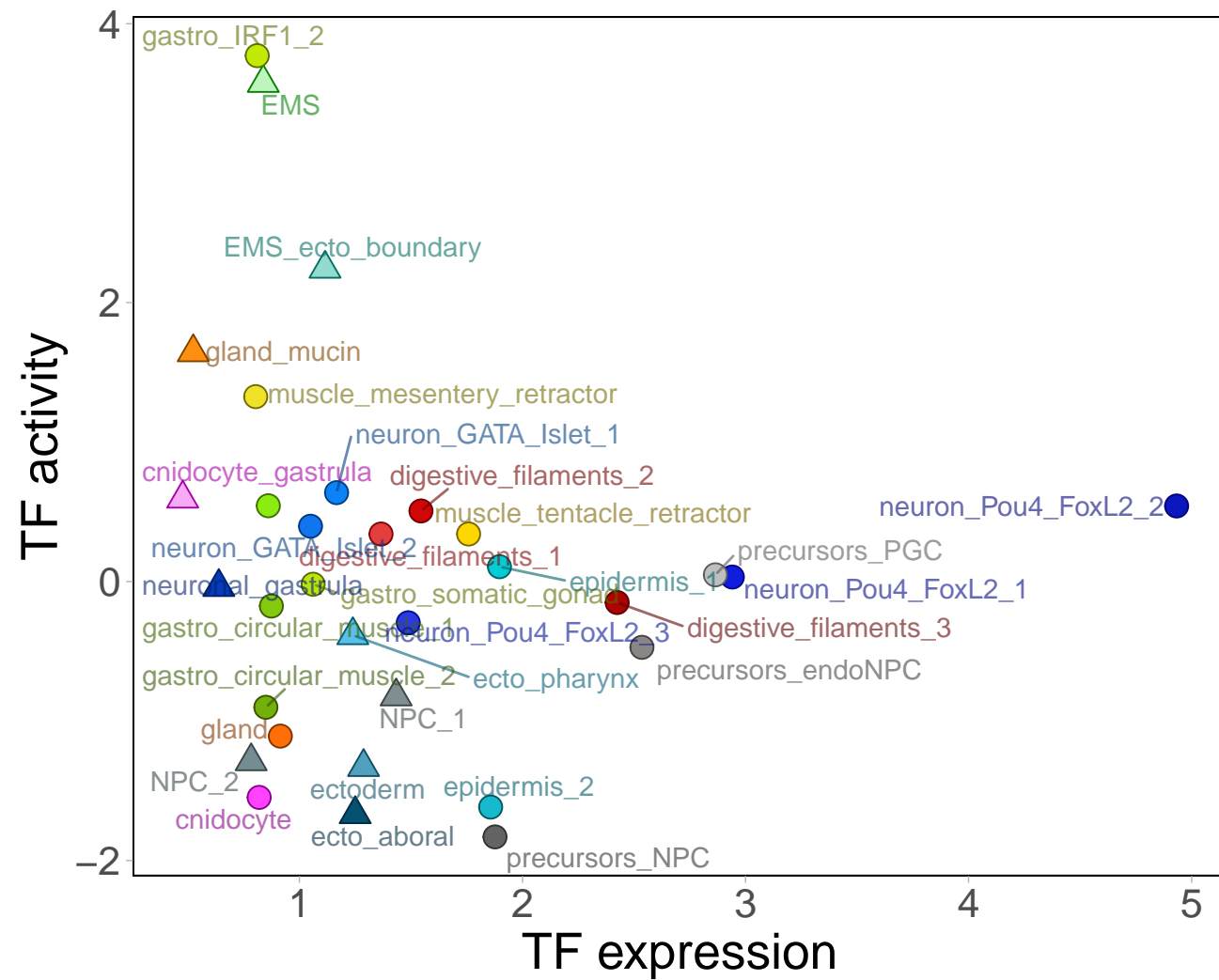
dimont



homer

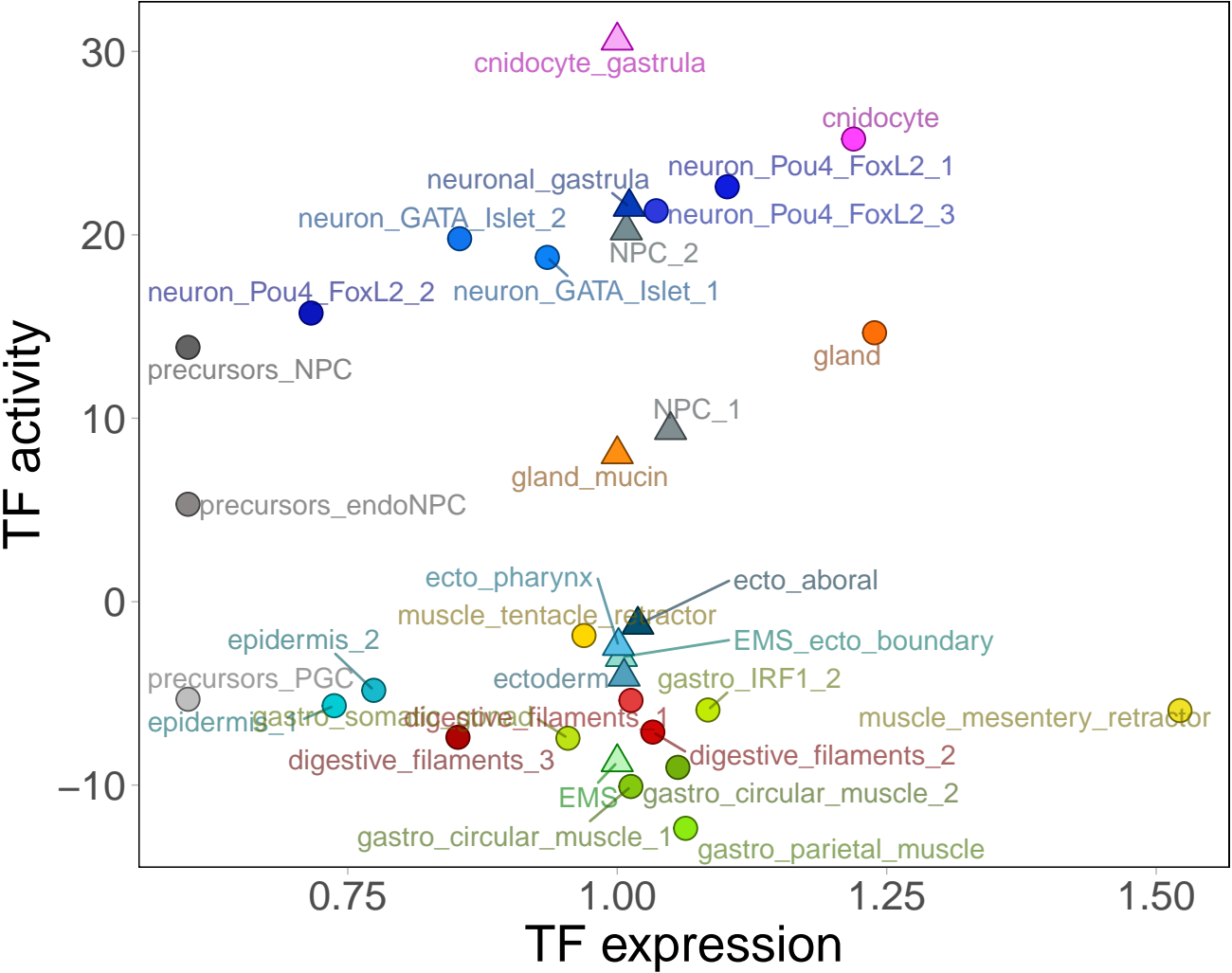


streme

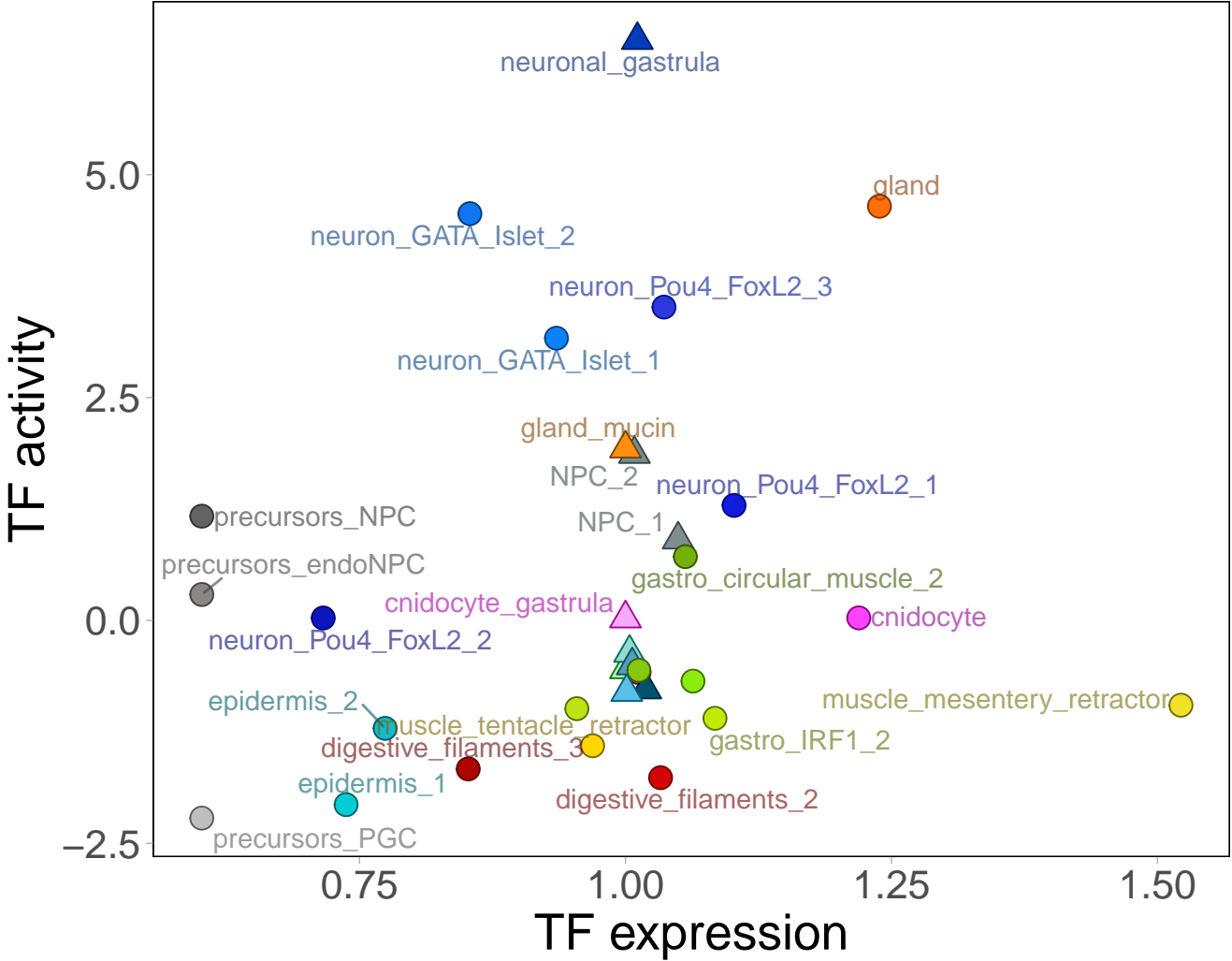


nvec_dap_34
Nvec_vc1.1_XM_032387318.2
OG: HLH.HG1.15:like:TFAP4
PFAM: HLH

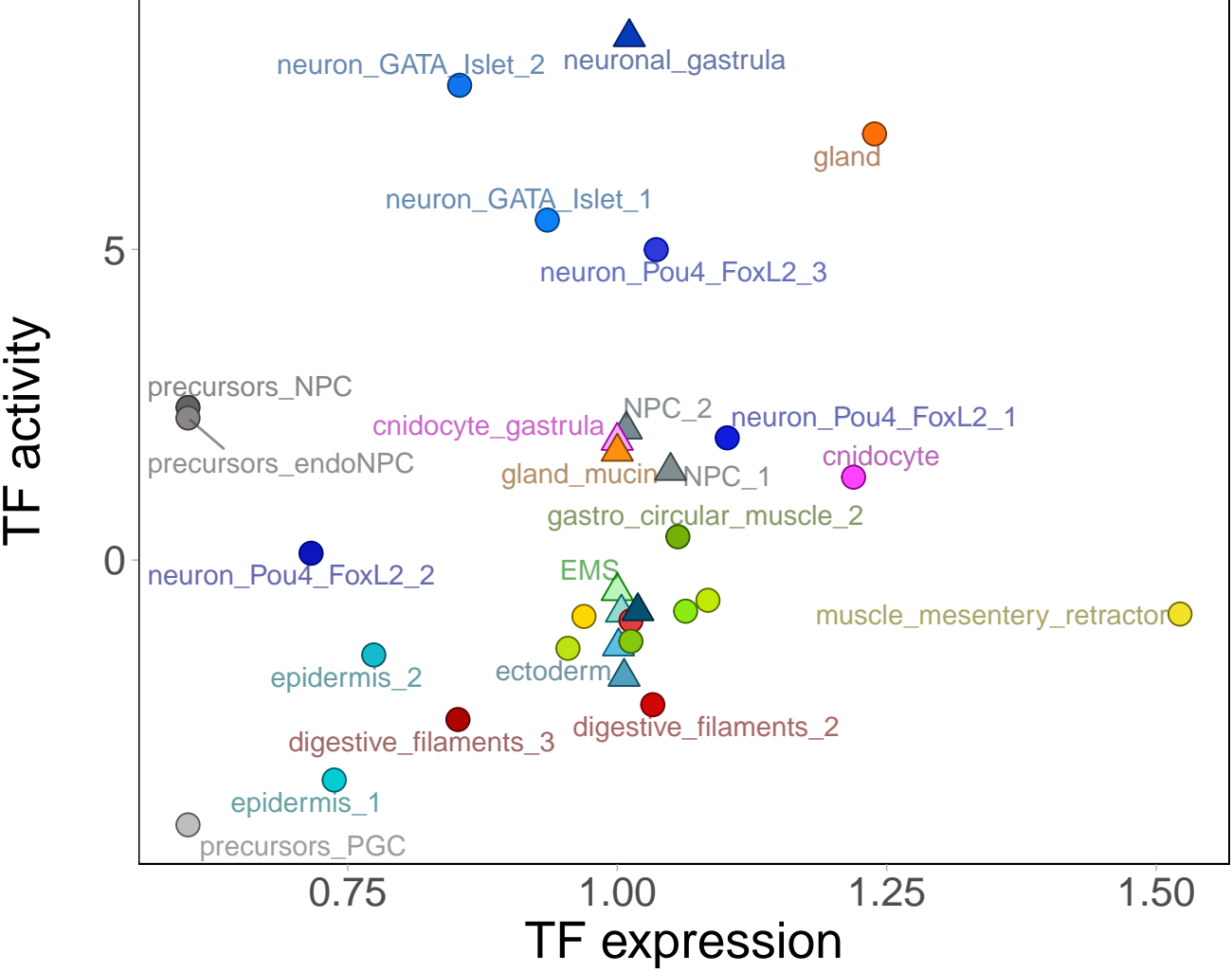
archetype



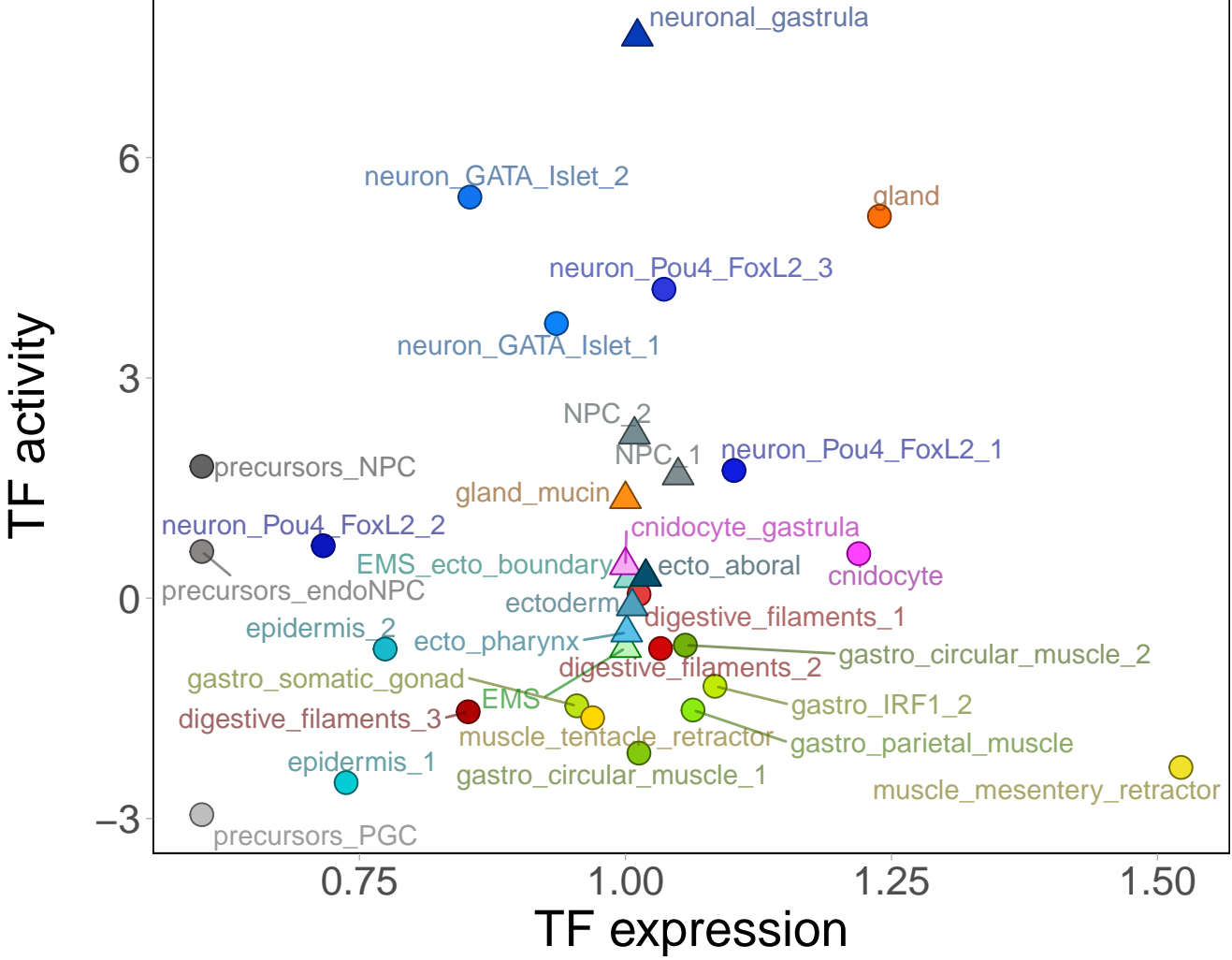
dimont



homer

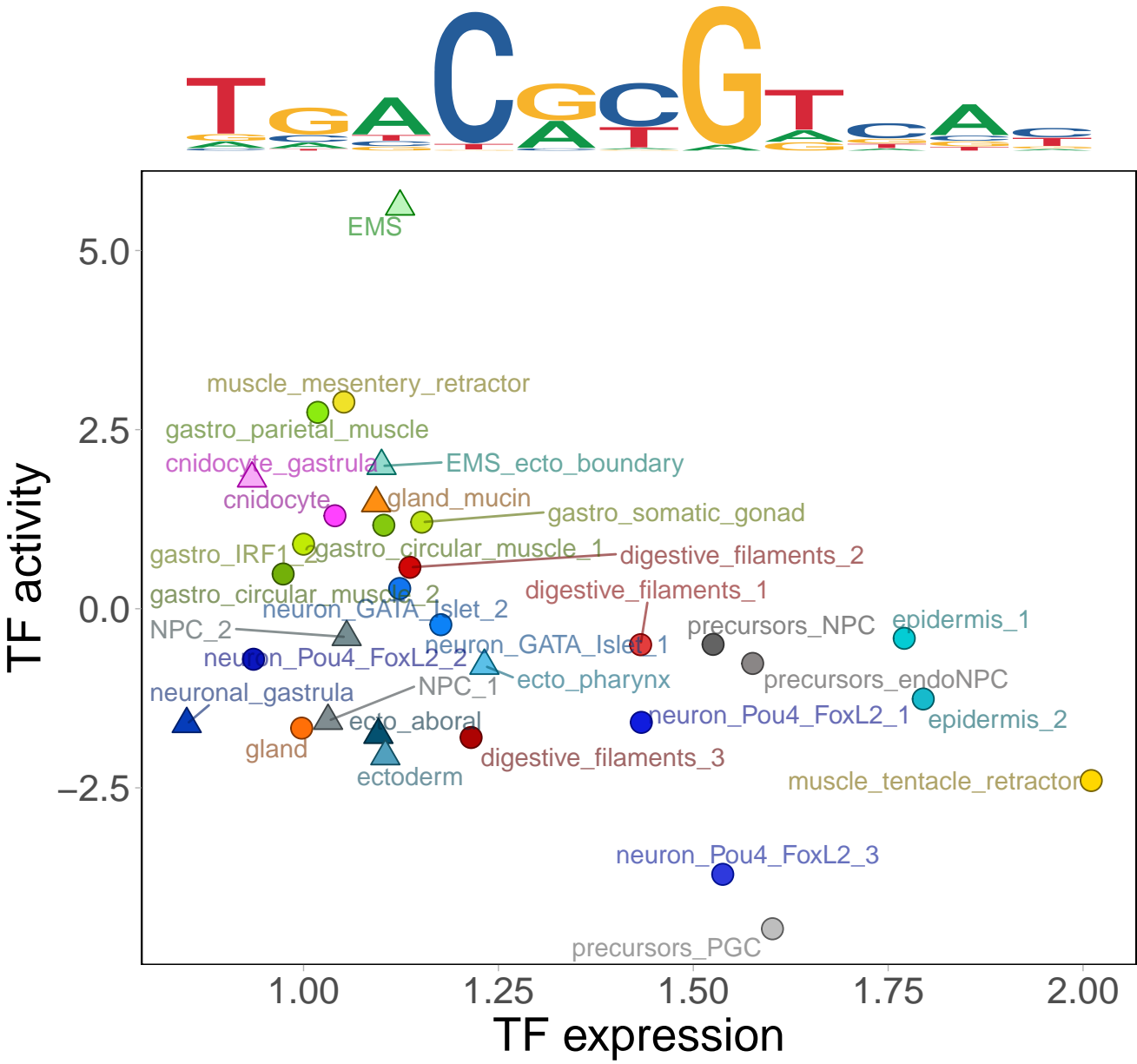


streme

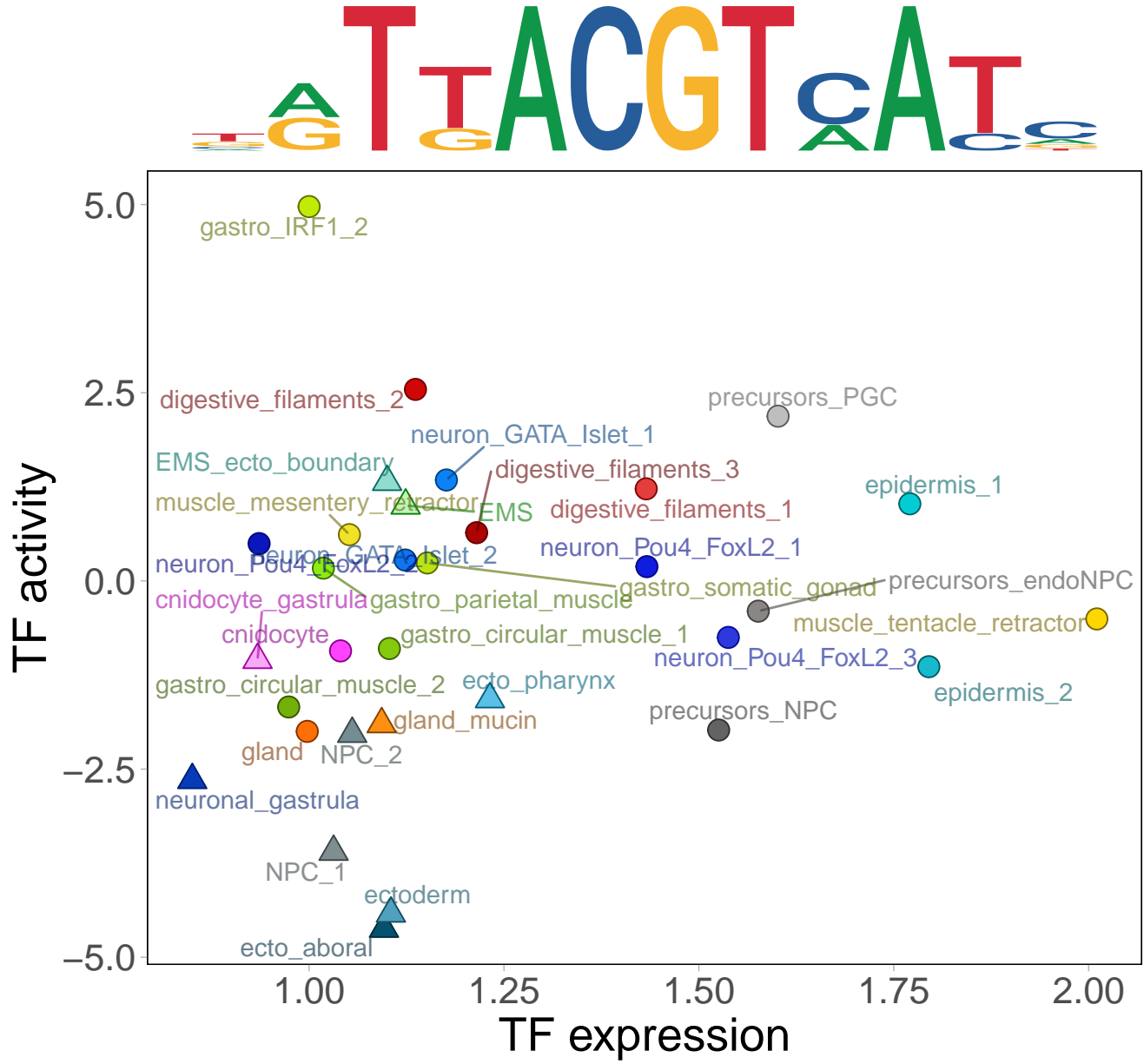


nvec_dap_35
Nvec_vc1.1_XM_032367127.2
OG: bZIP.HG2.10:DBP/HLF/TEF
PFAM: bZIP_2

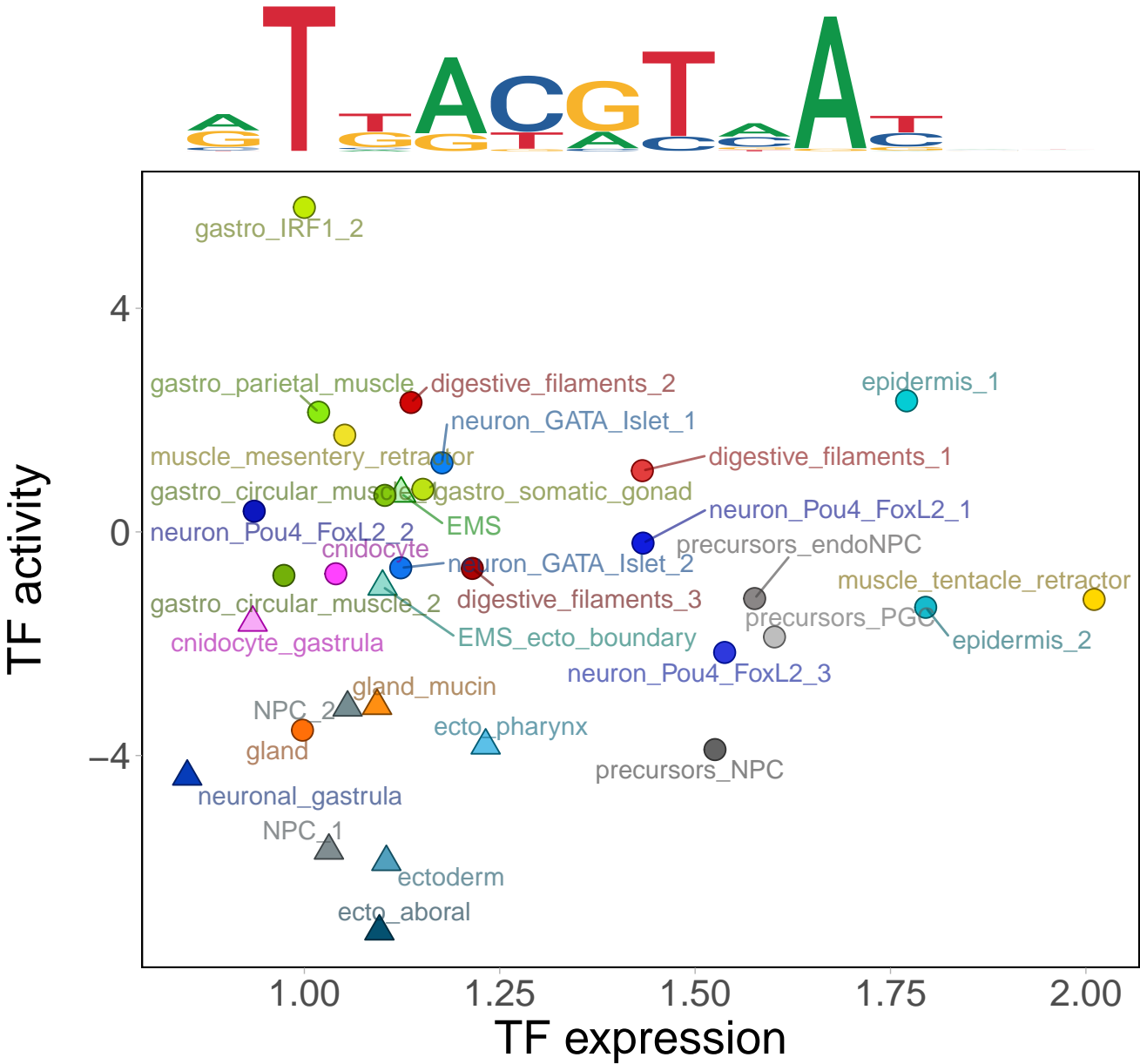
archetype



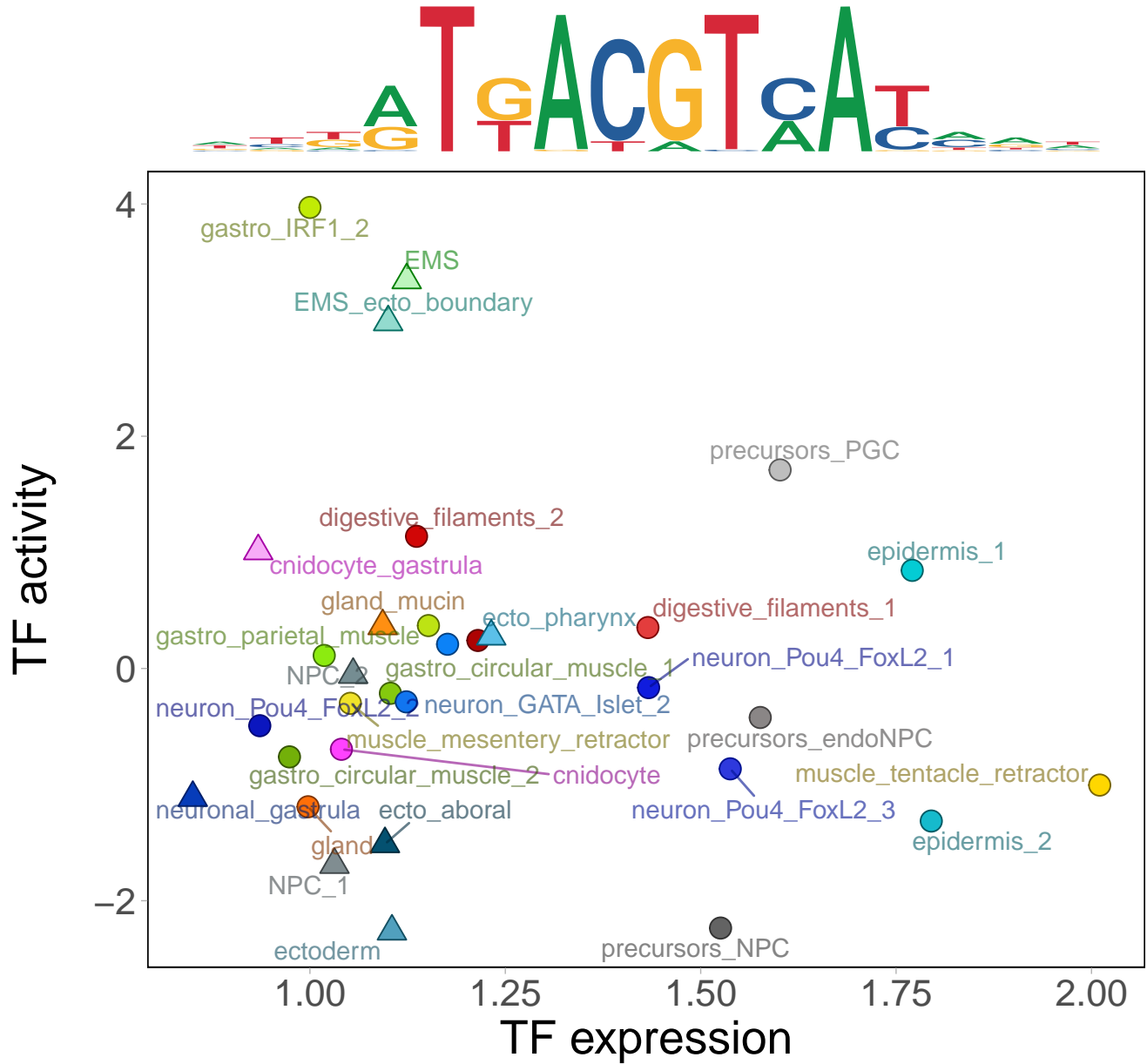
dimont



homer

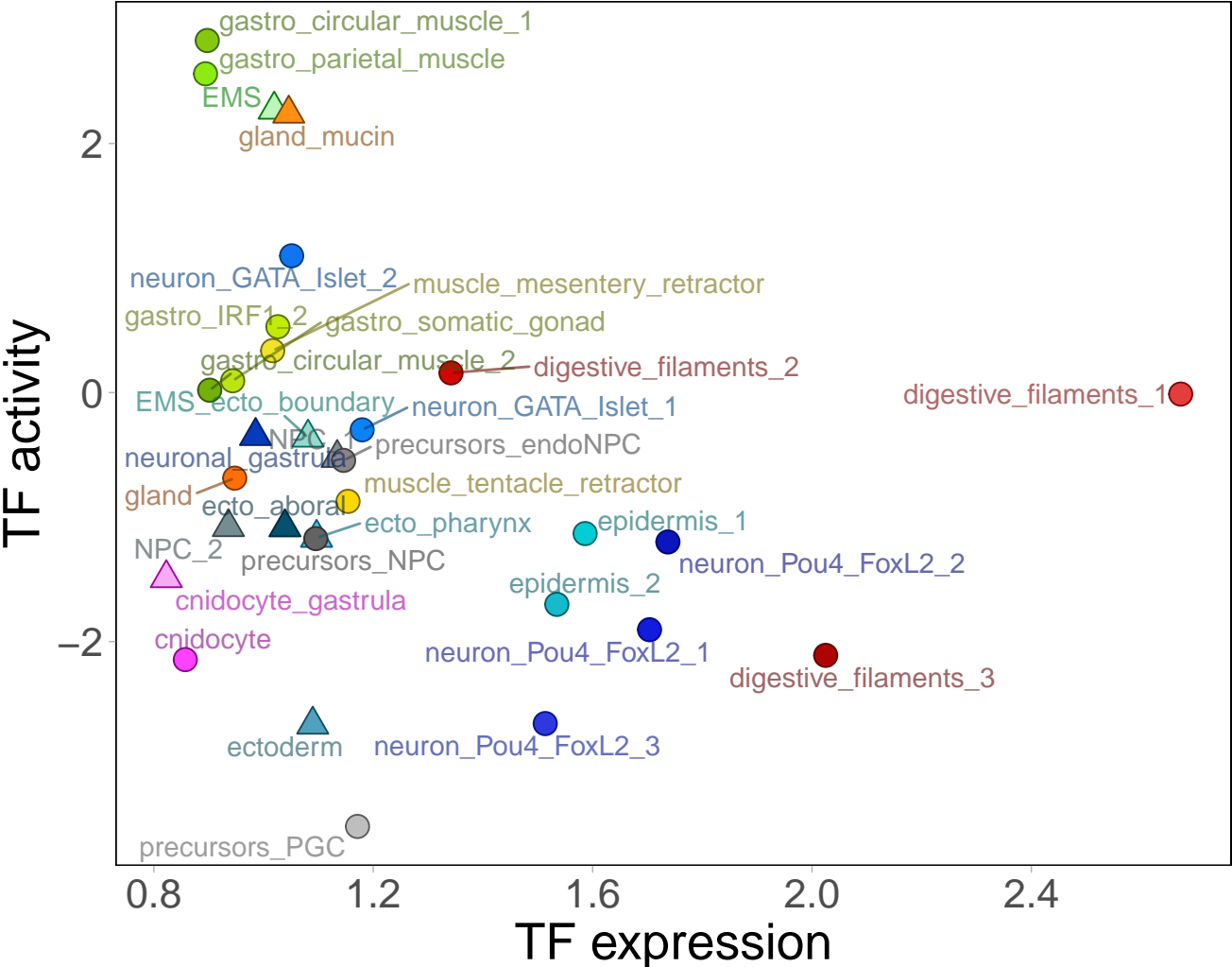


streame

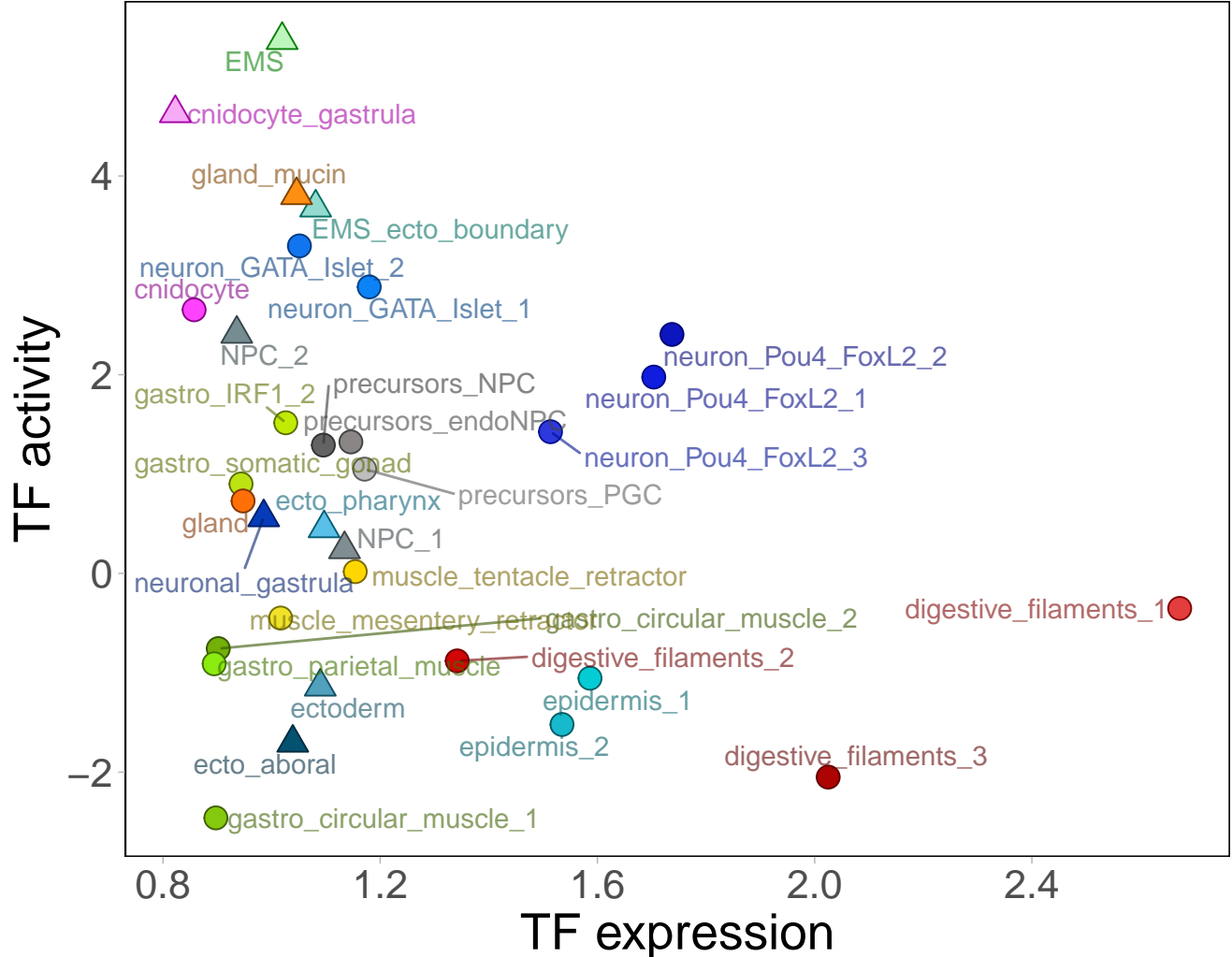


nvec_dap_36
Nvec_vc1.1_XM_001627049.3
OG: HLH.HG1.19:HELT
PFAM: HLH/Hairy_orange

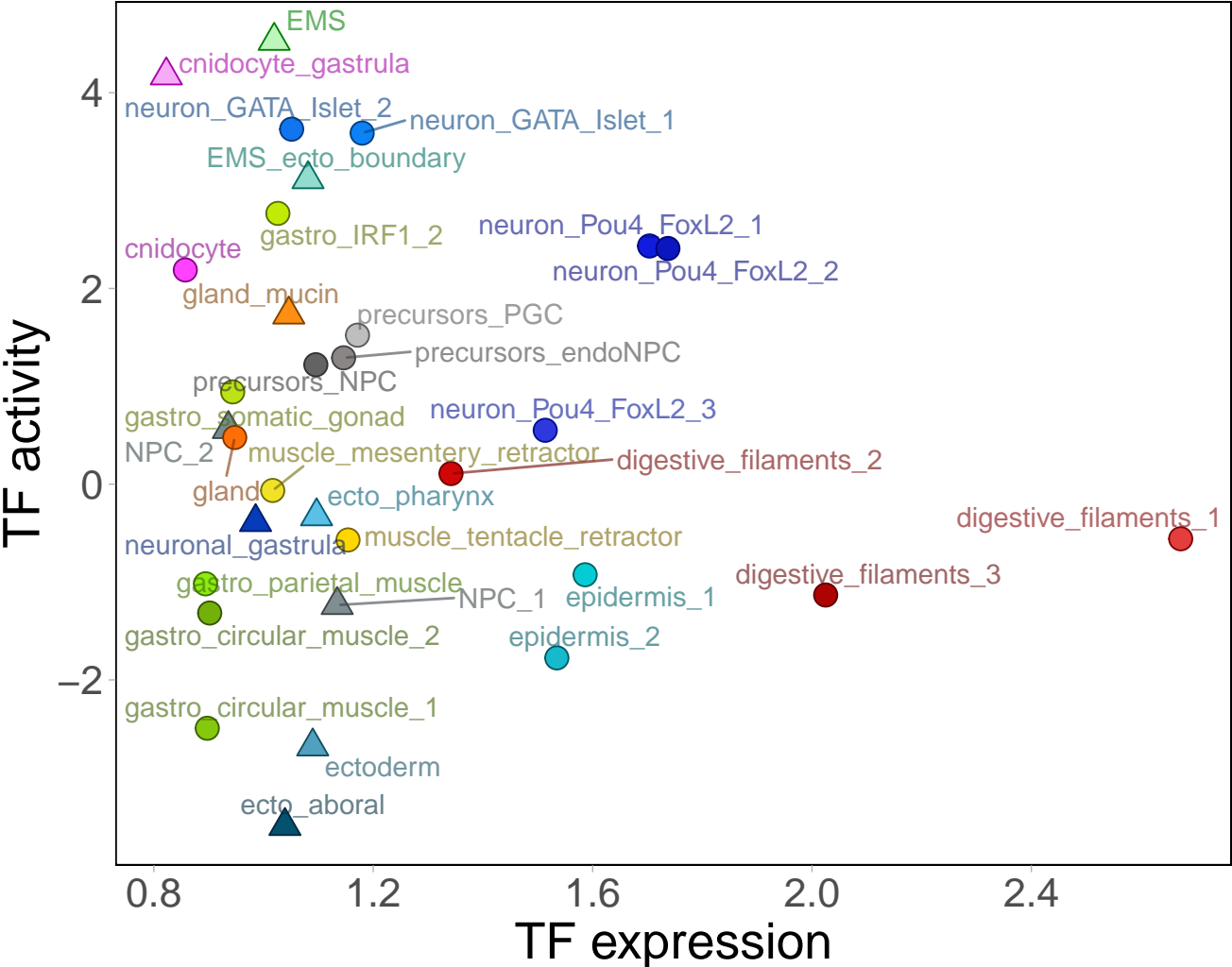
archetype



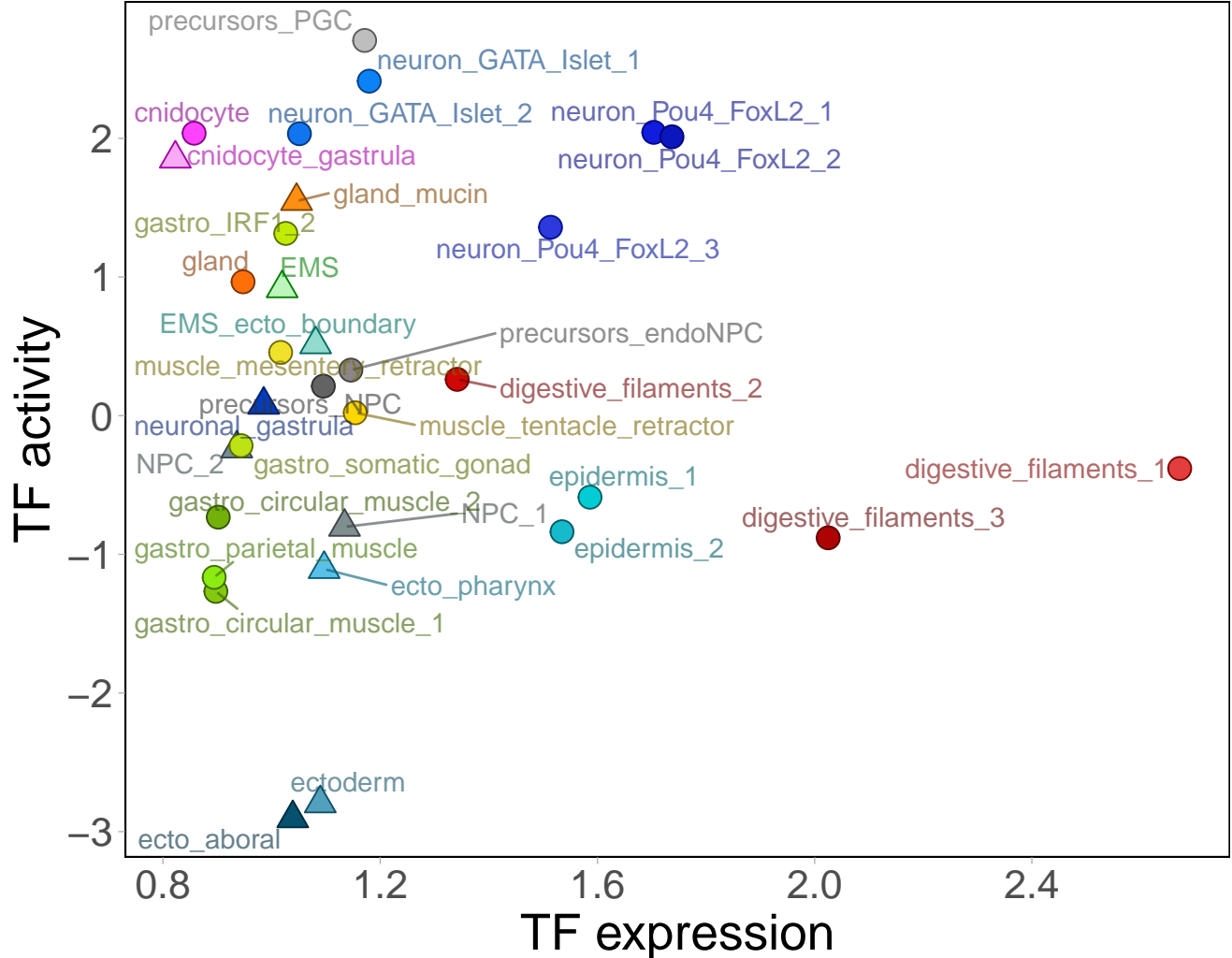
dimont



homer

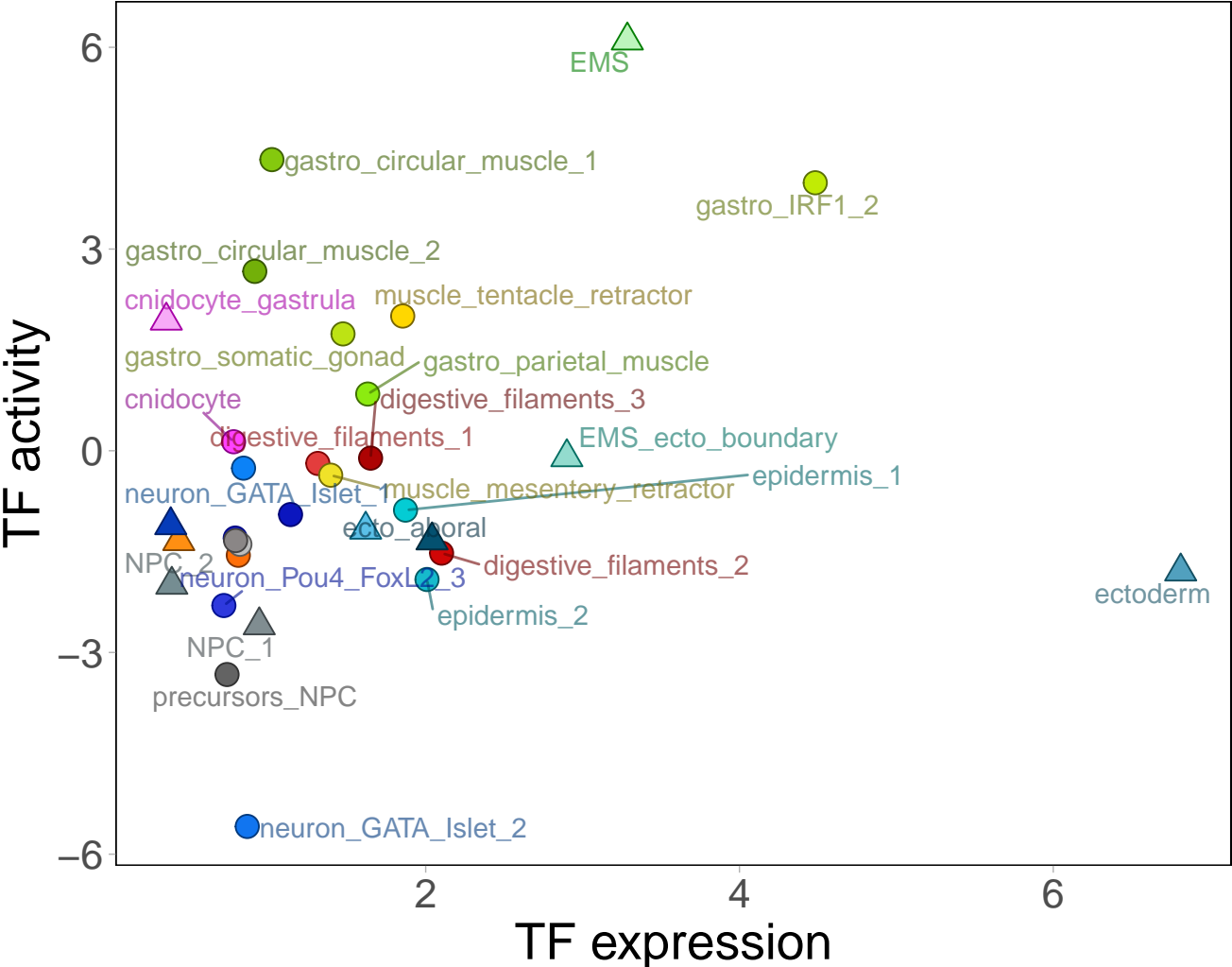


streame

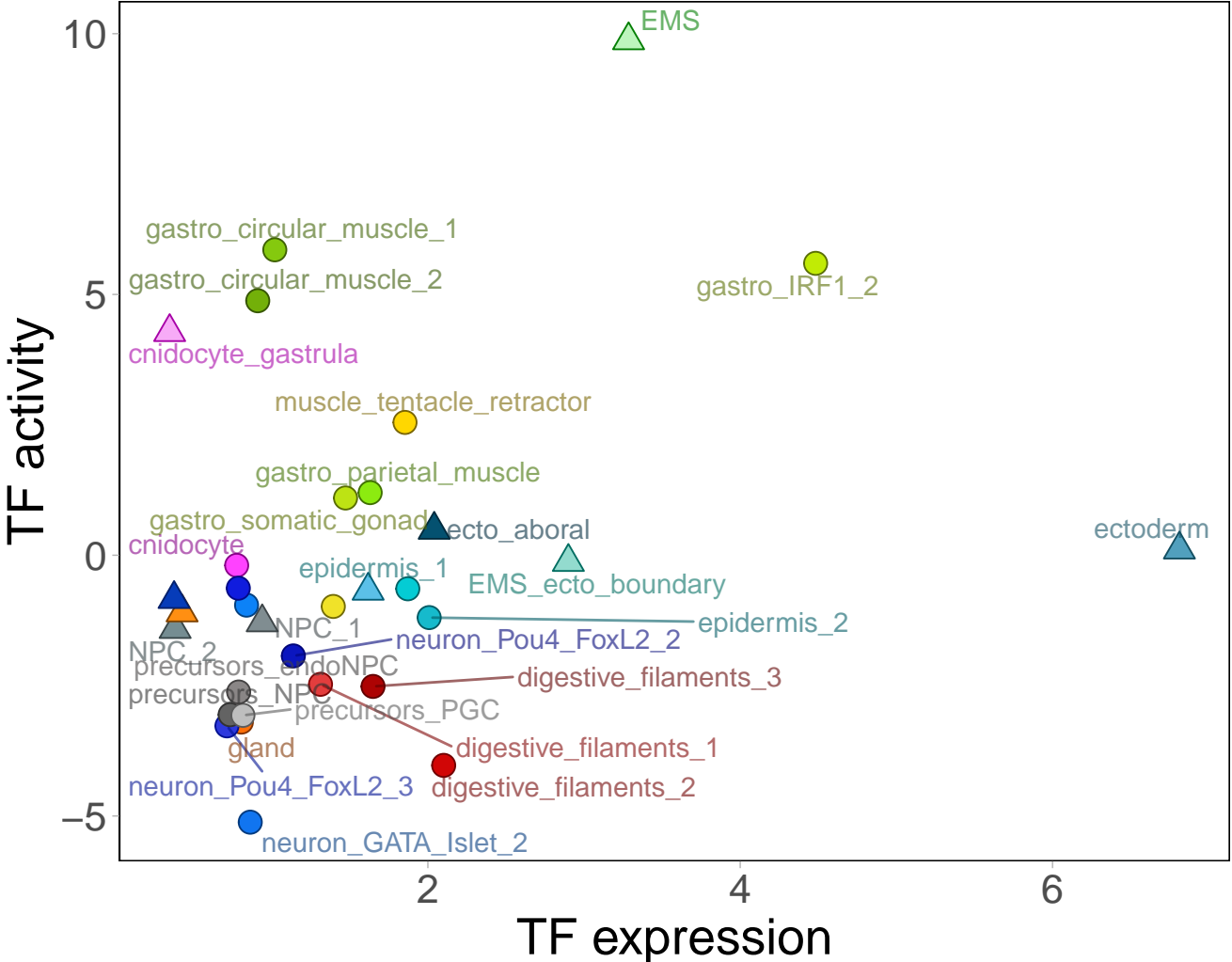


nvec_dap_37
Nvec_vc1.1_XM_032381586.2
OG: zf-C2H2.HG2.21:KLF5
PFAM: zf-C2H2/zf-C2H2/zf-C2H2

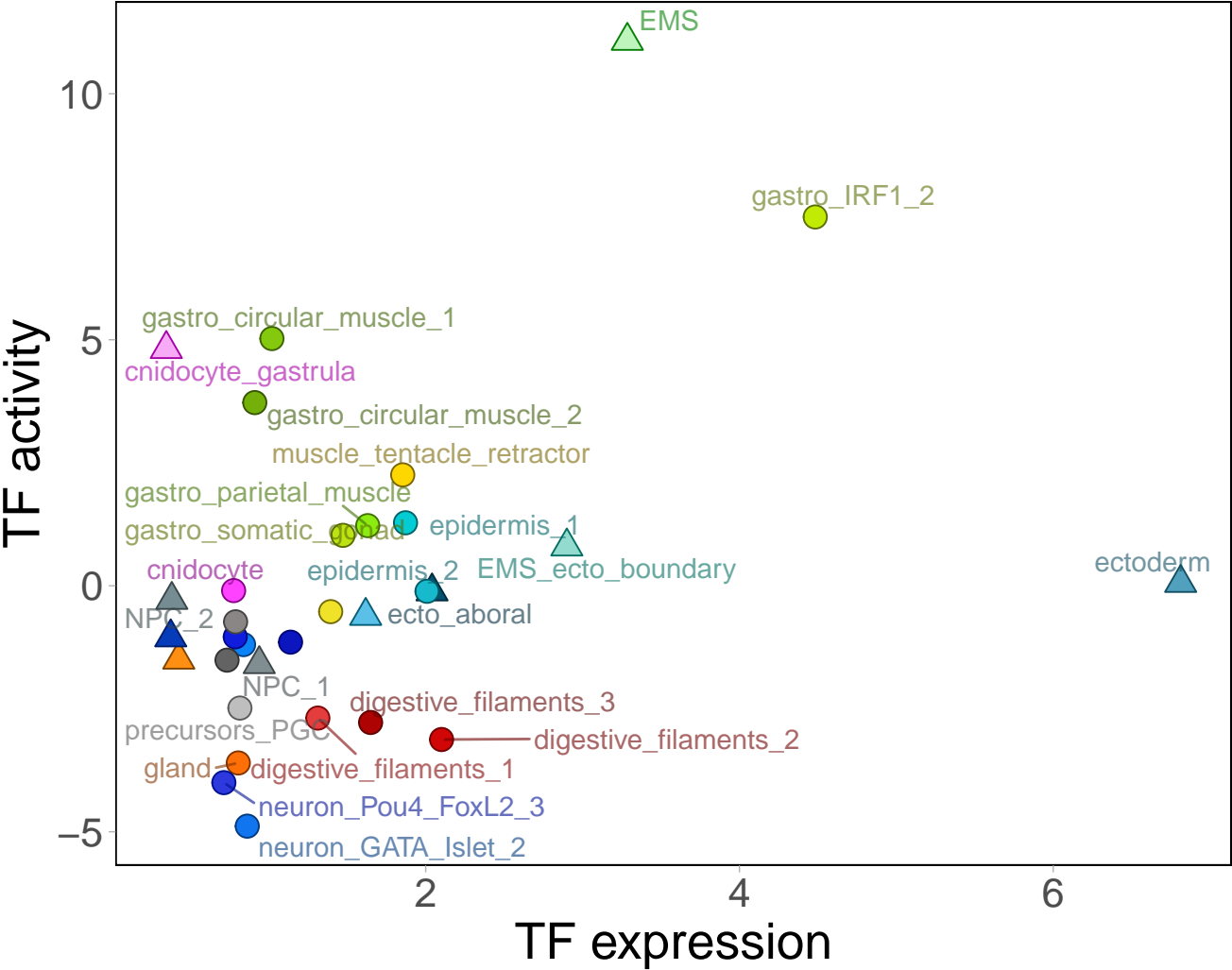
archetype



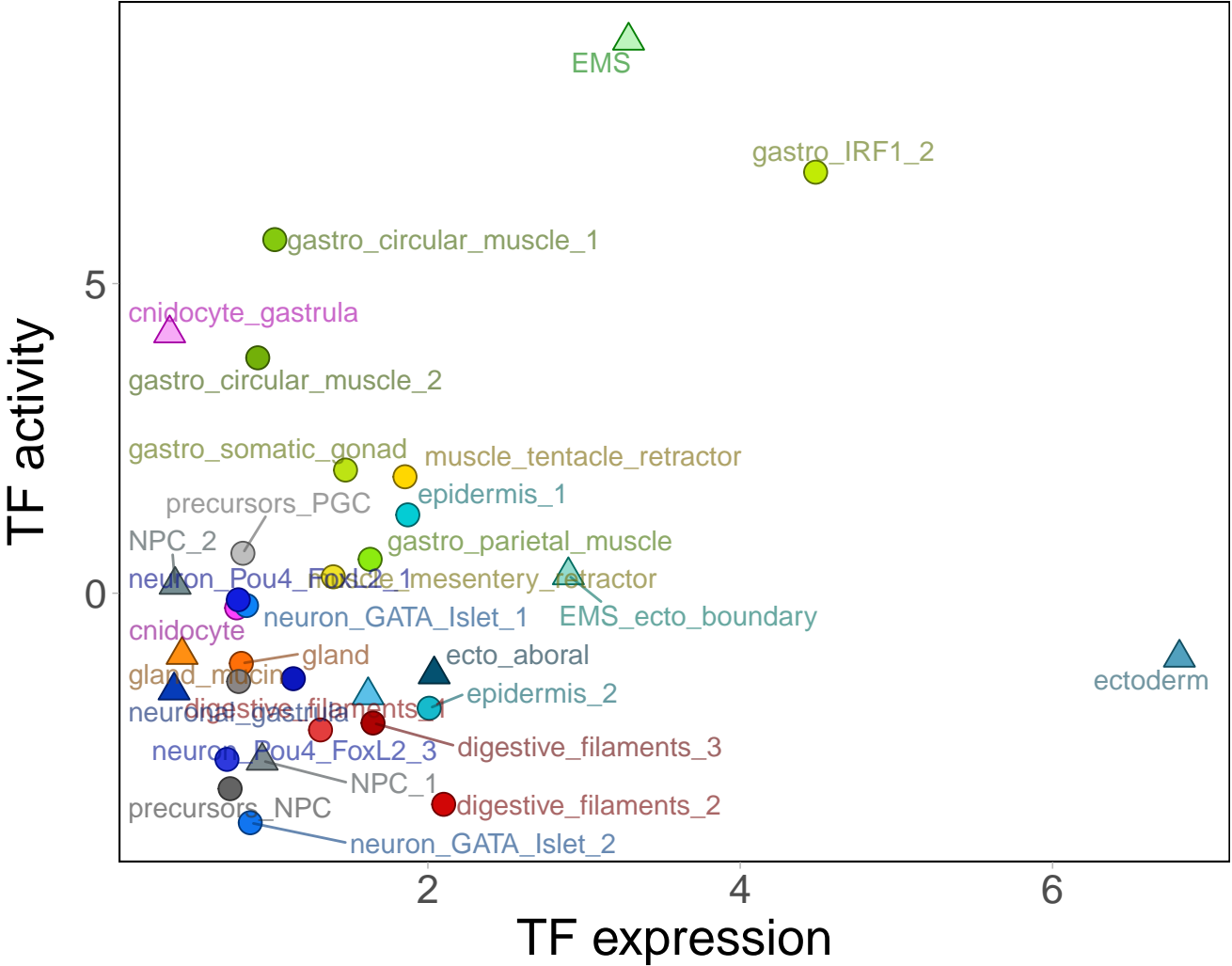
dimont



homer

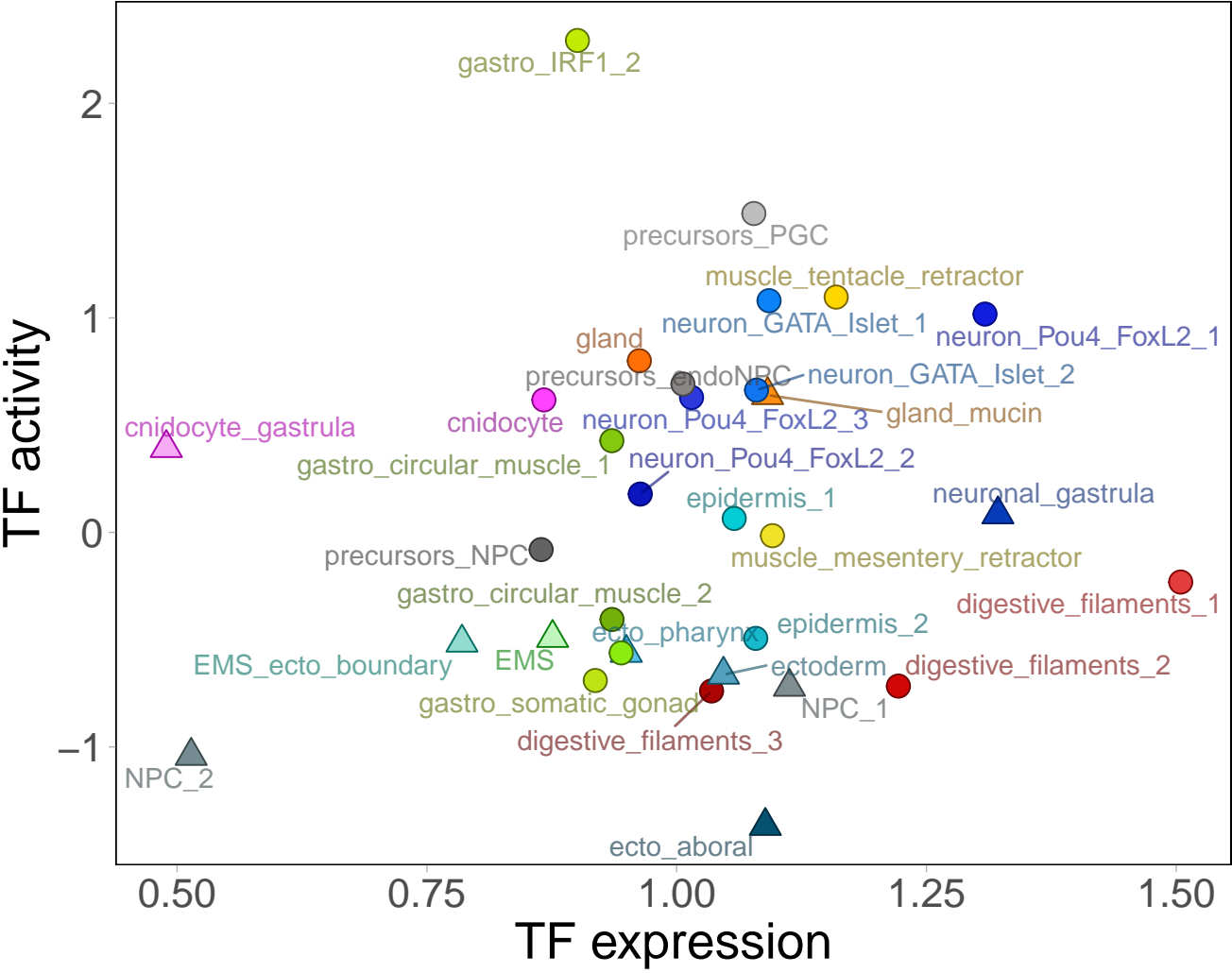


streame

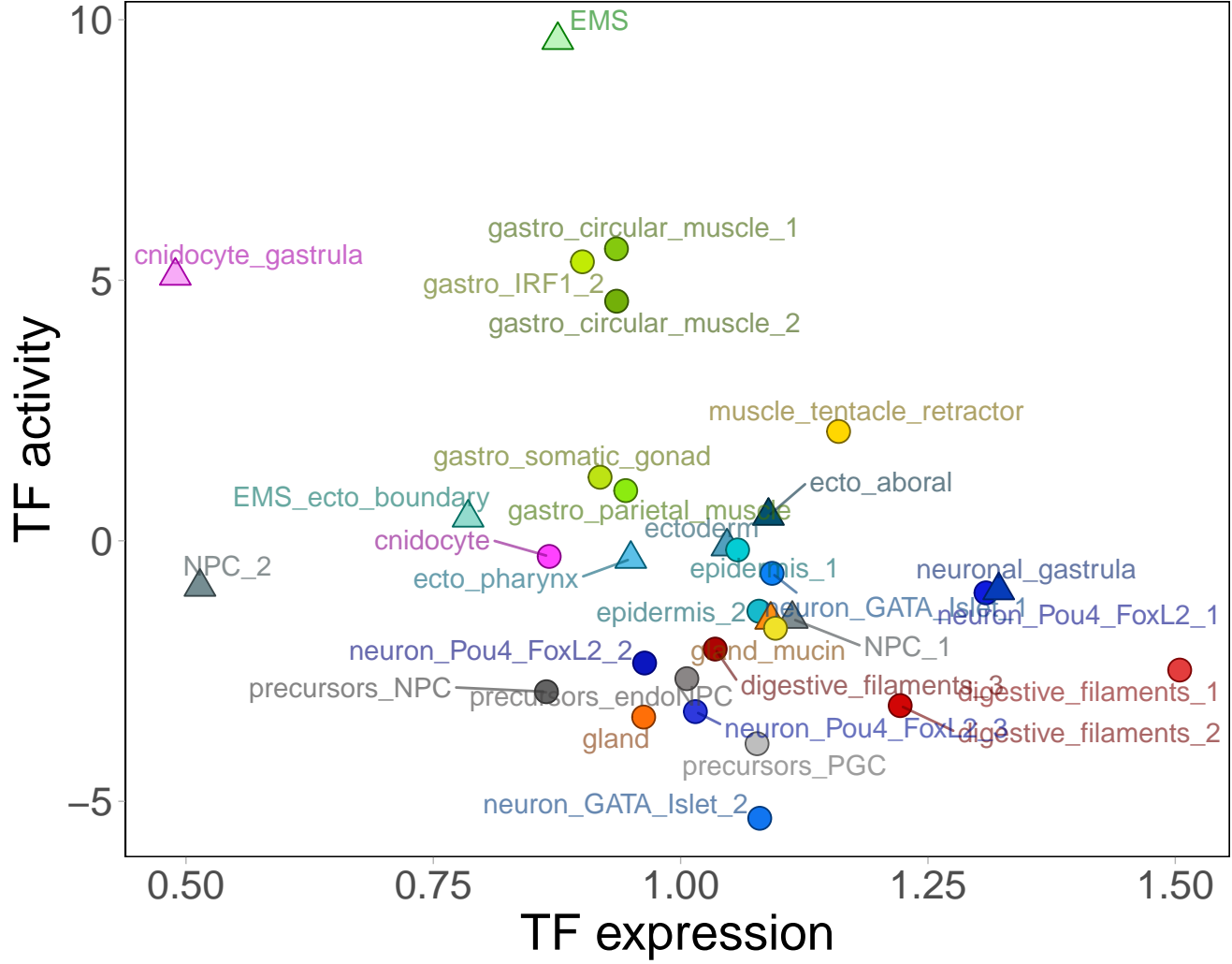


nvec_dap_39
Nvec_vc1.1_XM_032381562.2
OG: zf-C2H2.HG2.18:like:KLF5/KLF6/KLF7
PFAM: zf-C2H2/zf-C2H2/zf-C2H2

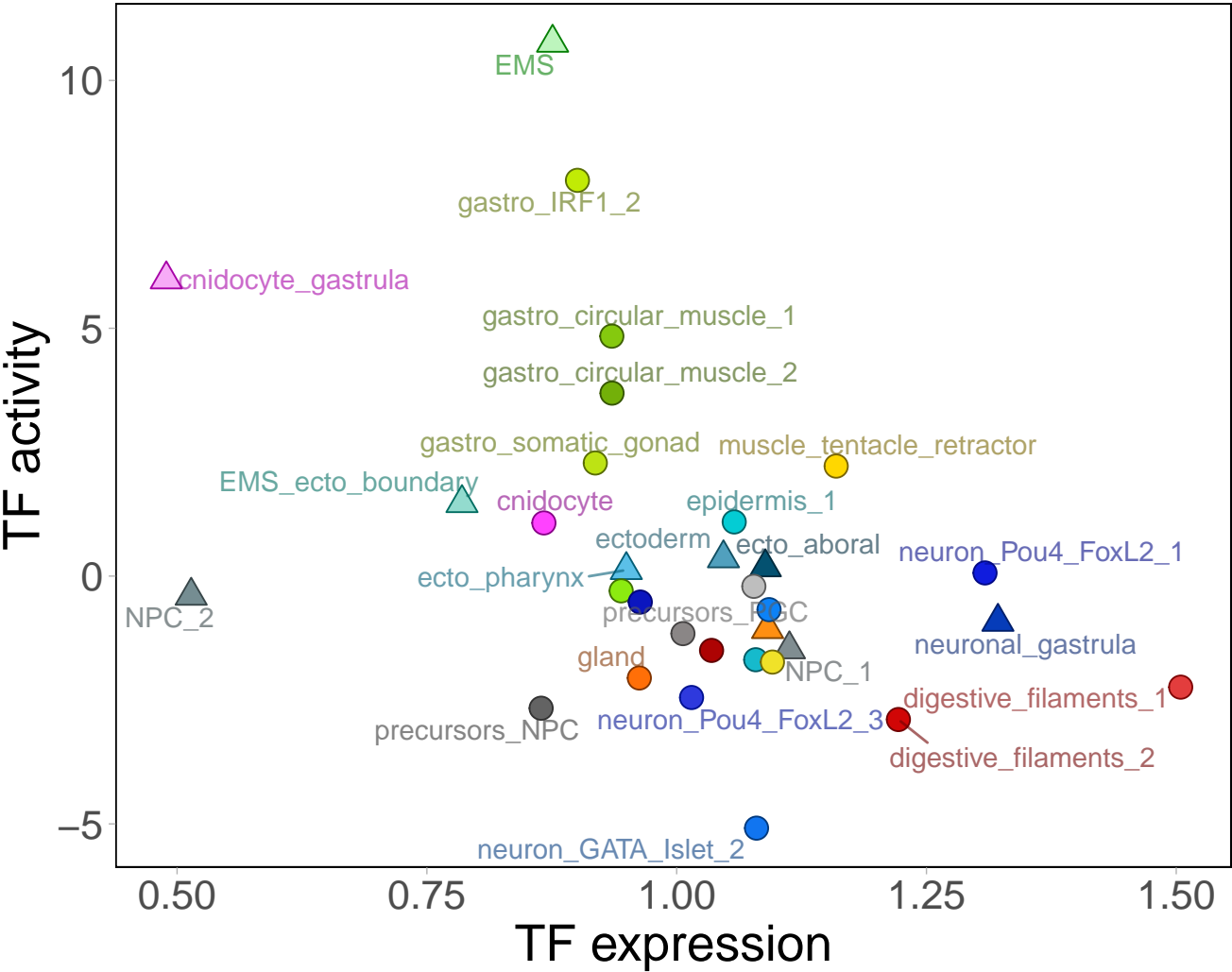
archetype



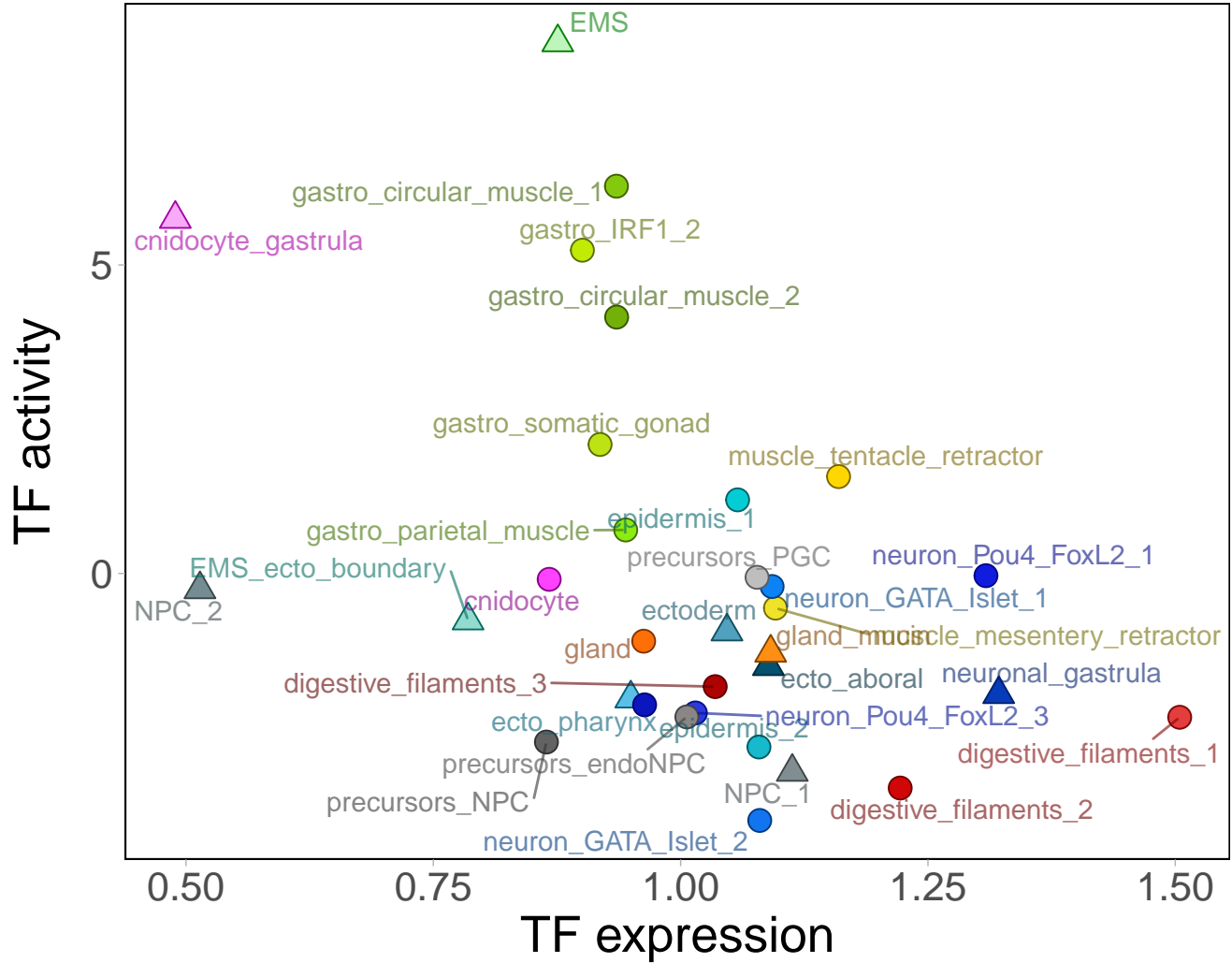
dimont



homer

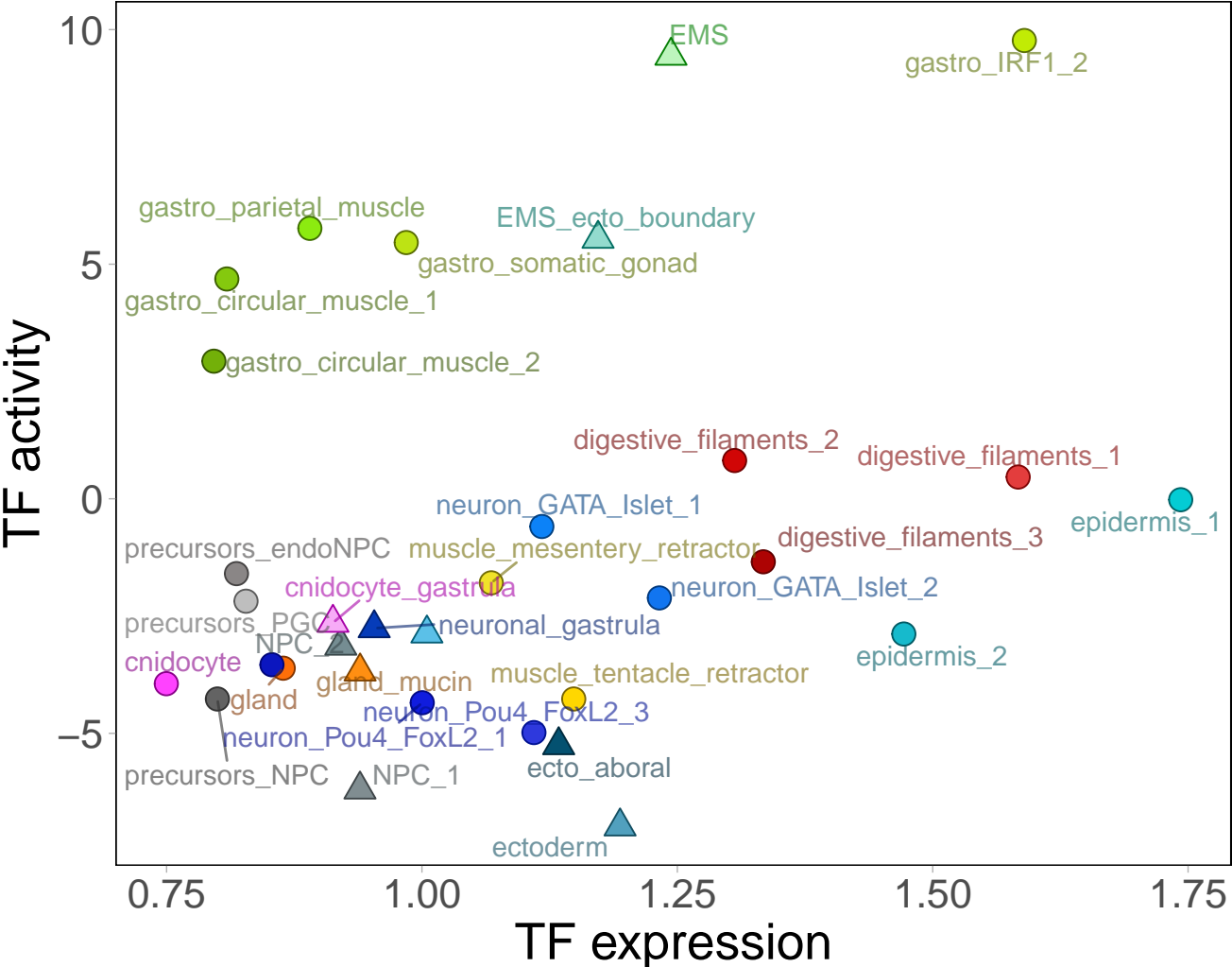


streame

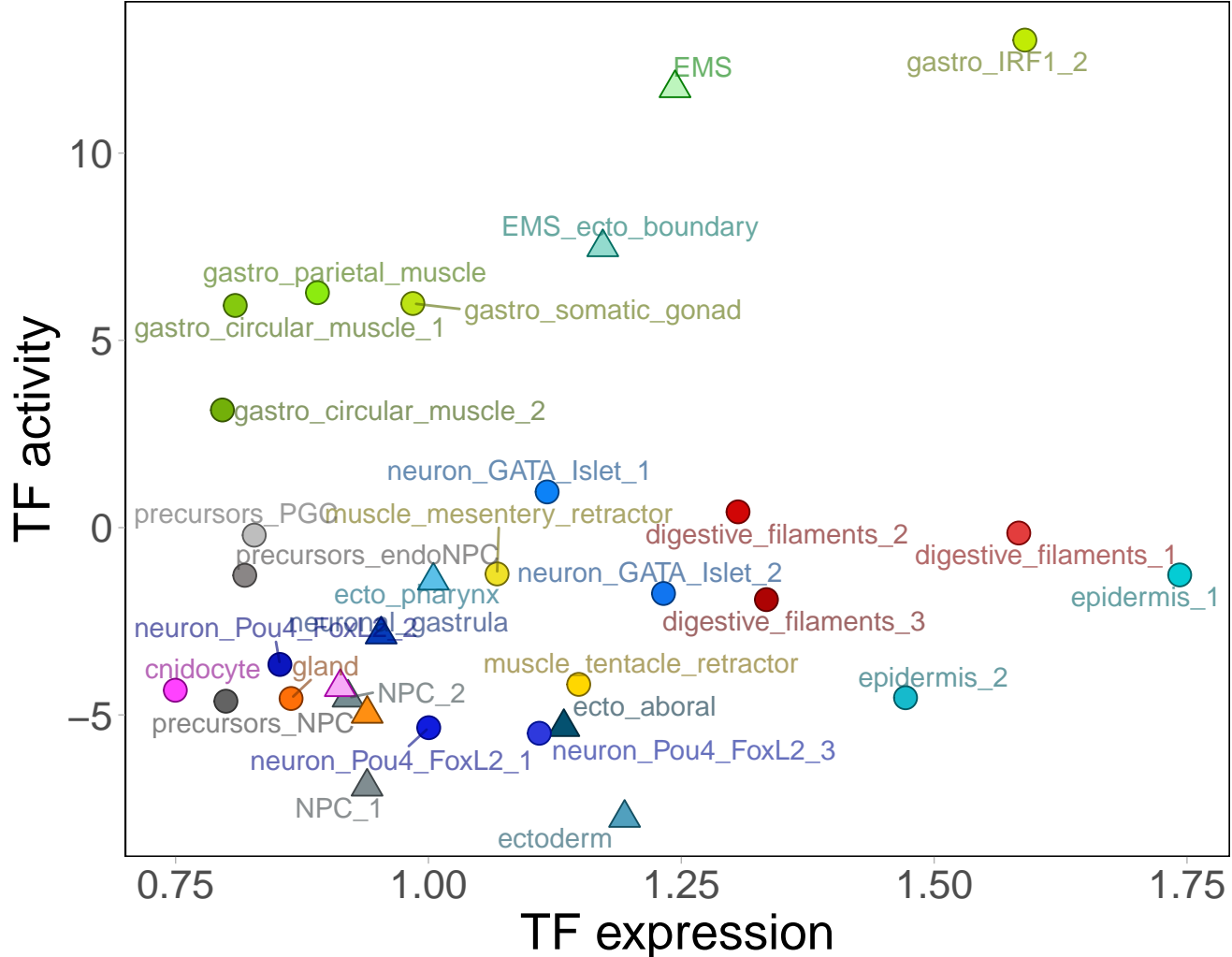


nvec_dap_40
Nvec_vc1.1_XM_048732532.1
OG: Ets.HG1.8:ETS1/ETS2
PFAM: Ets

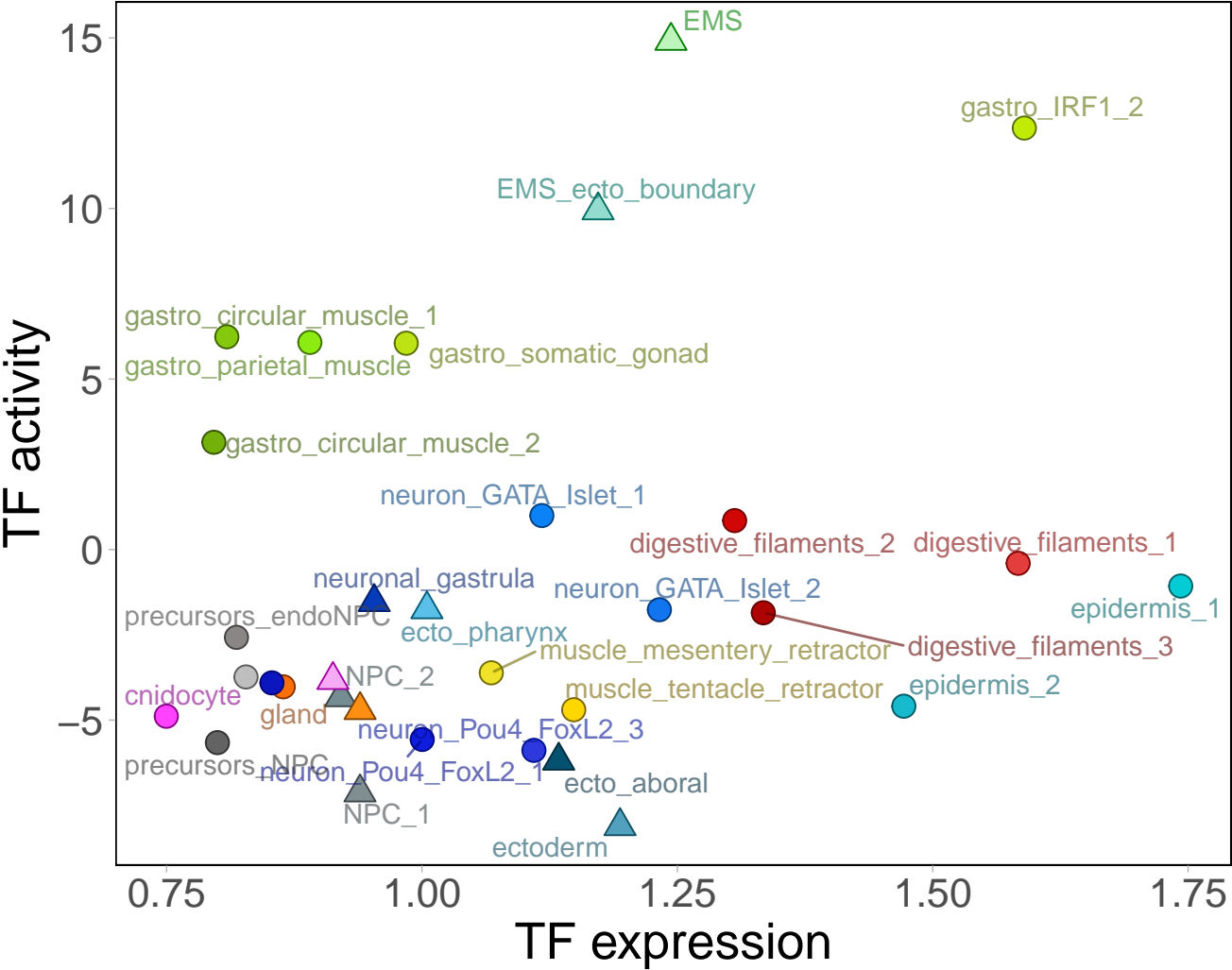
archetype



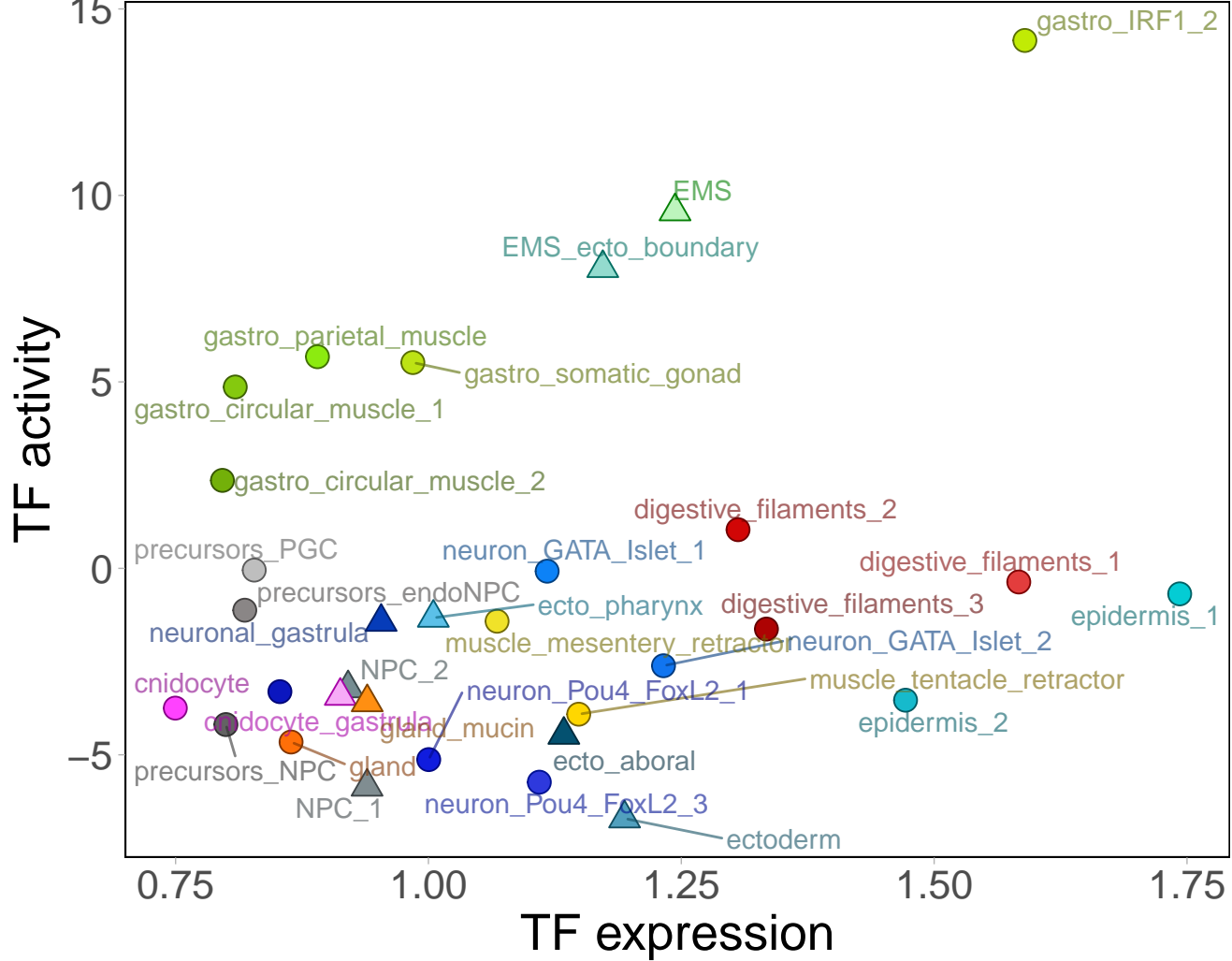
dimont



homer

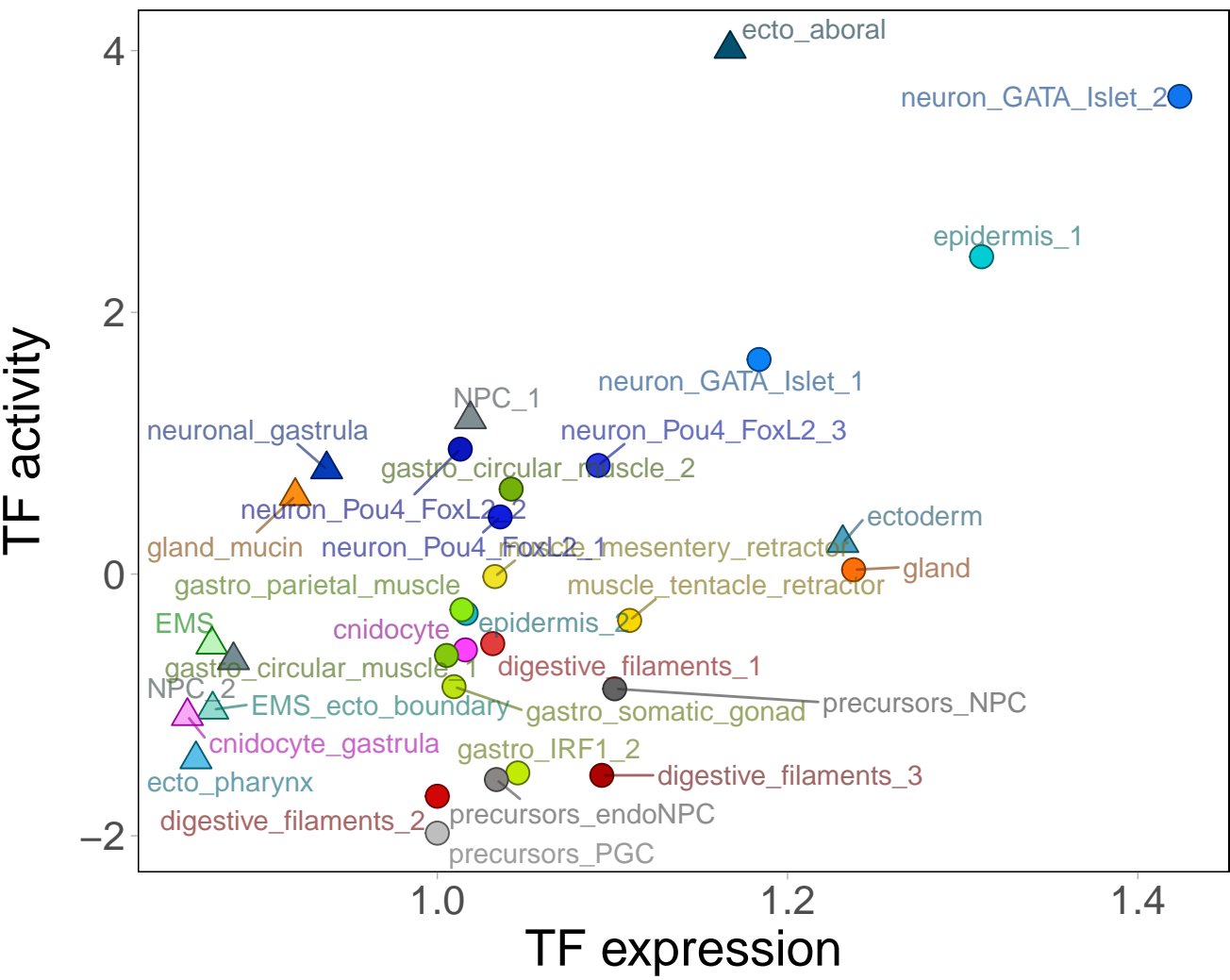


streame

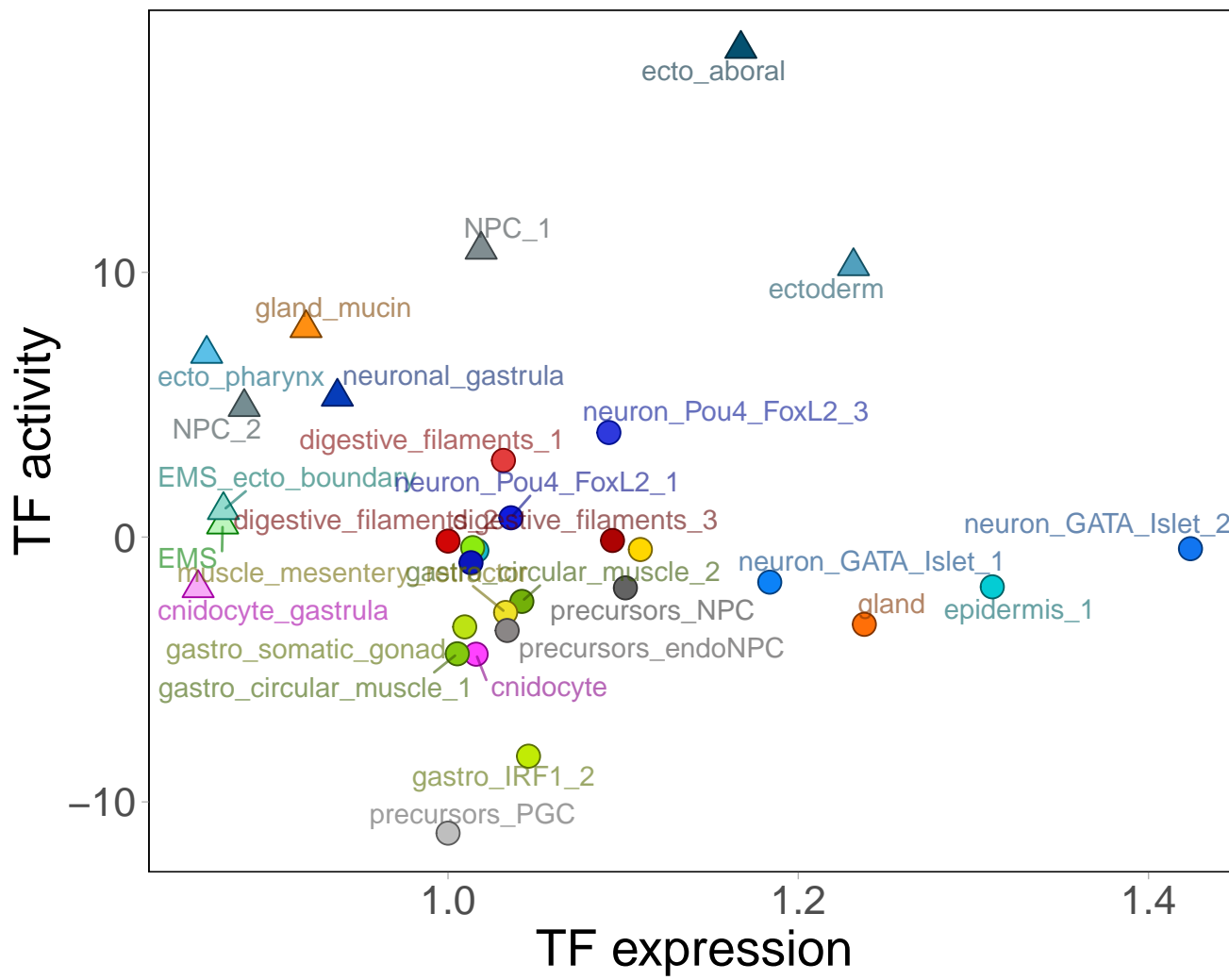


nvec_dap_41
Nvec_vc1.1_XM_001634160.3
OG: Homeodomains.HG2.57:RAX/RAX2
PFAM: Homeodomain/OAR

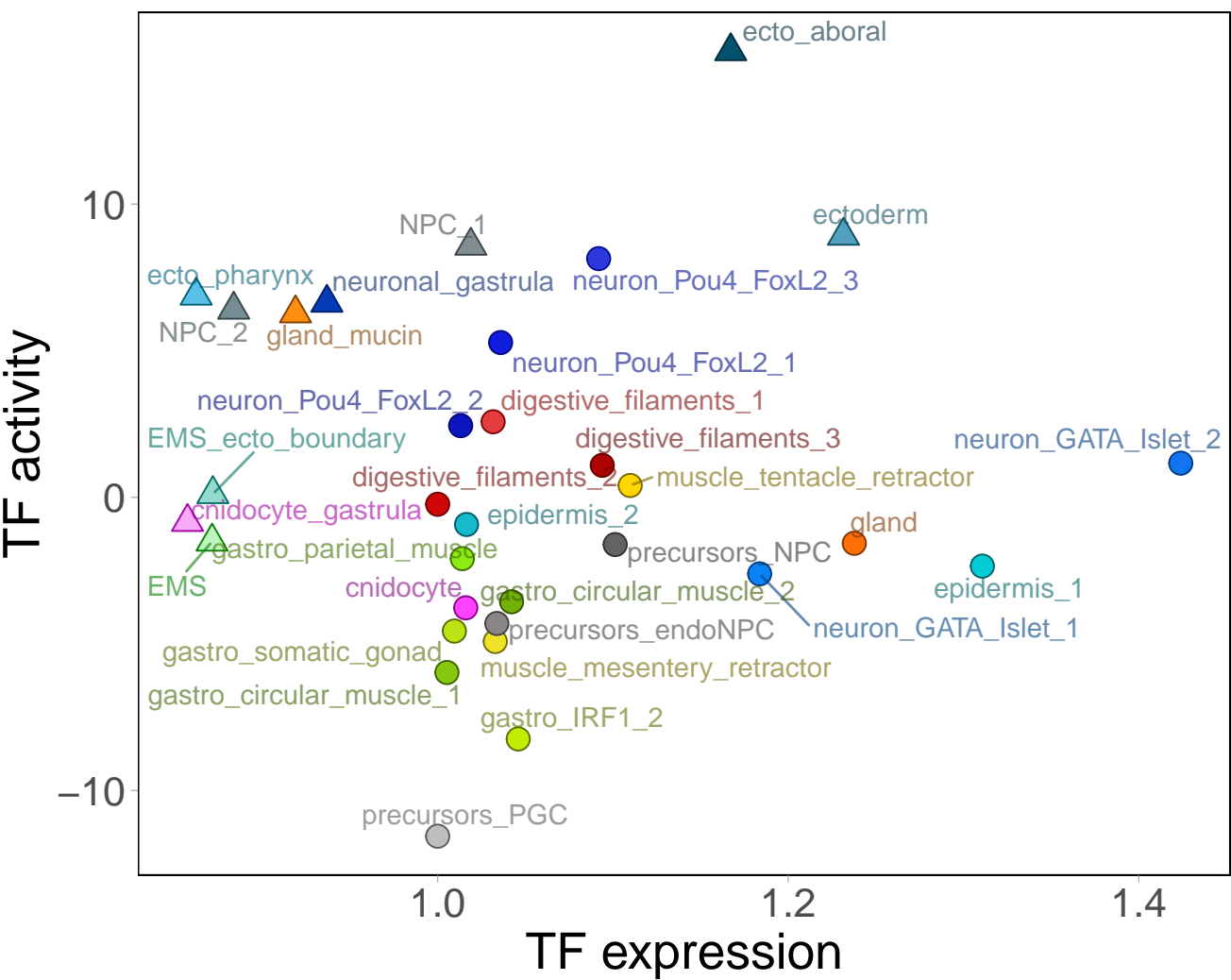
archetype



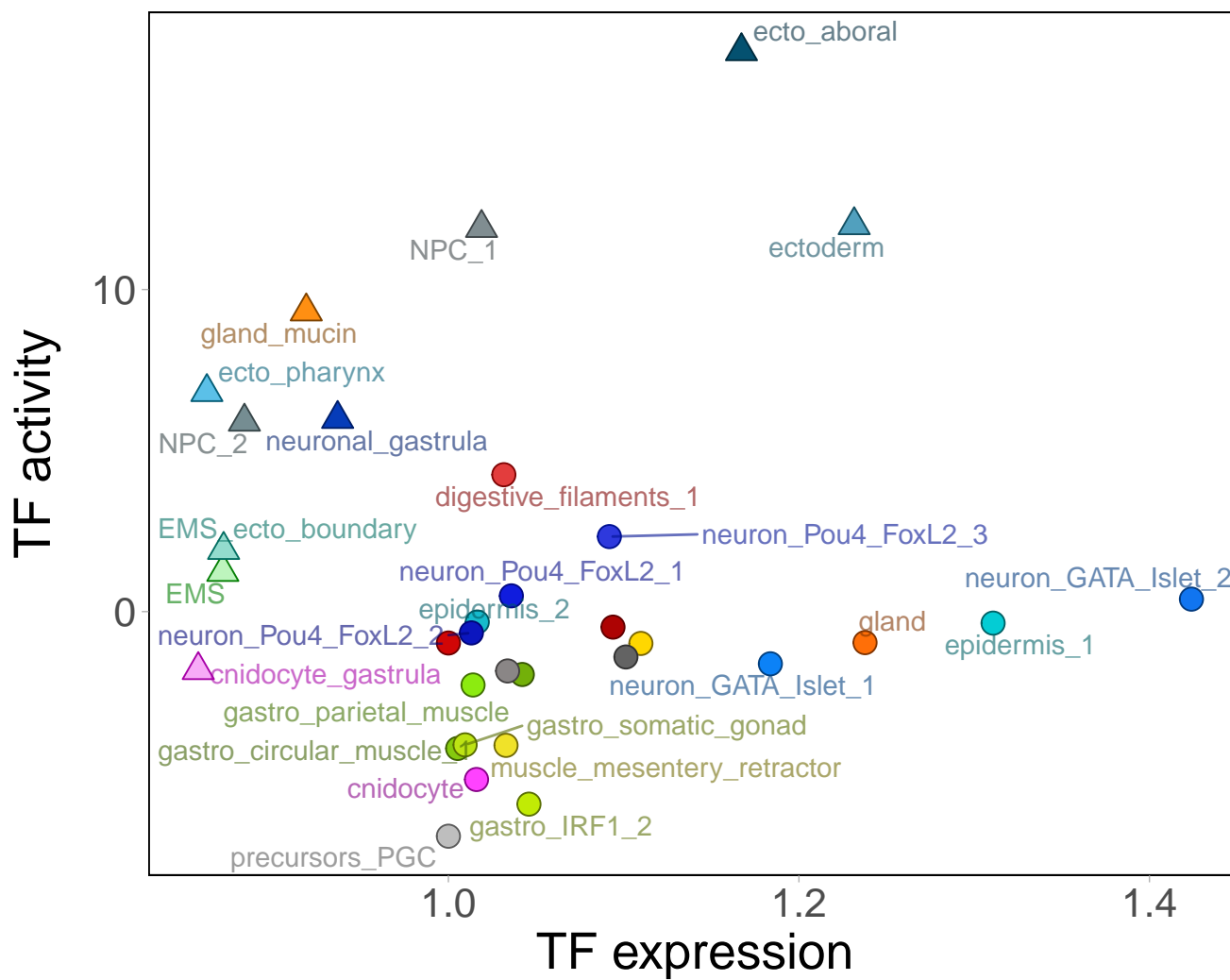
dimont



homer

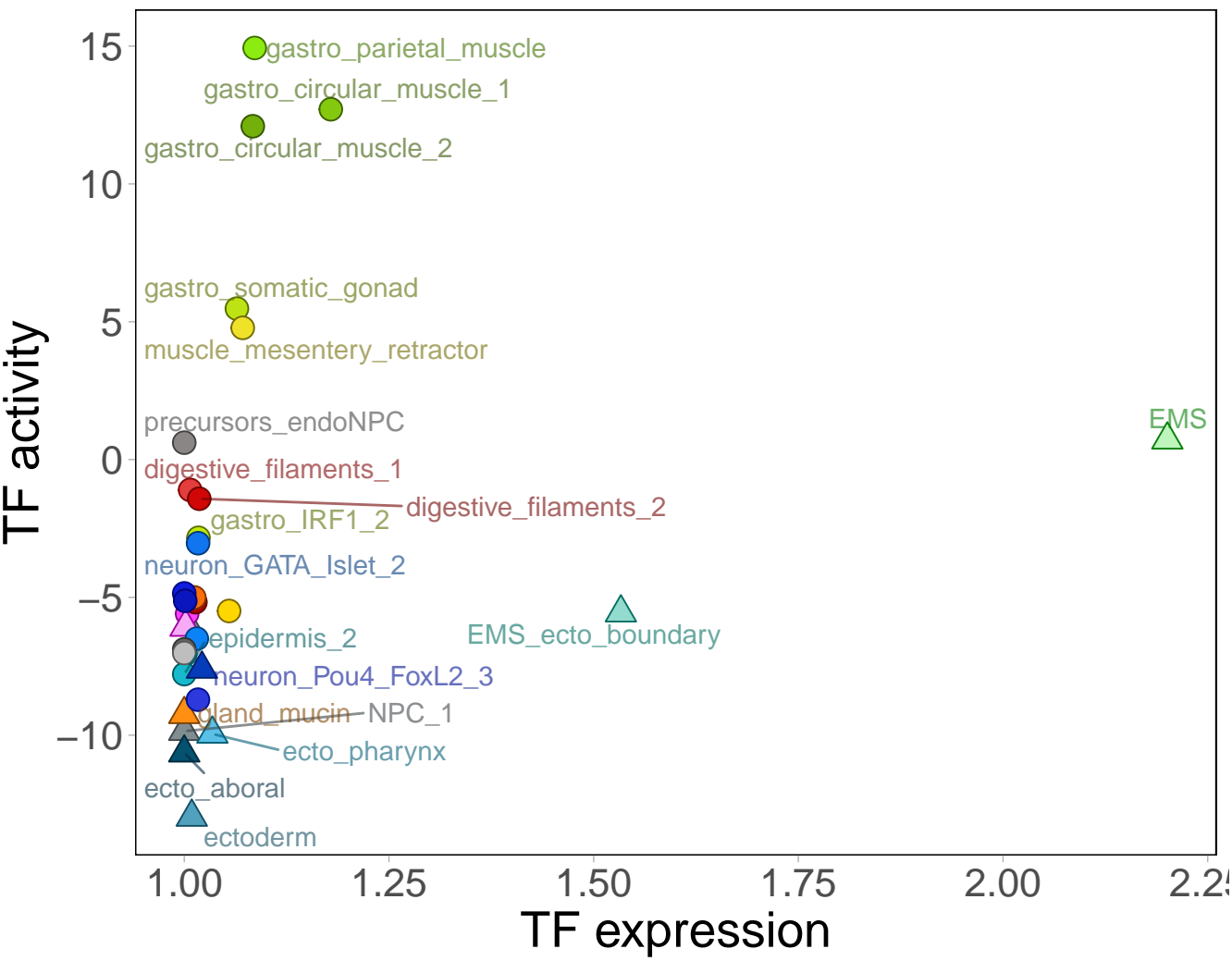


streame

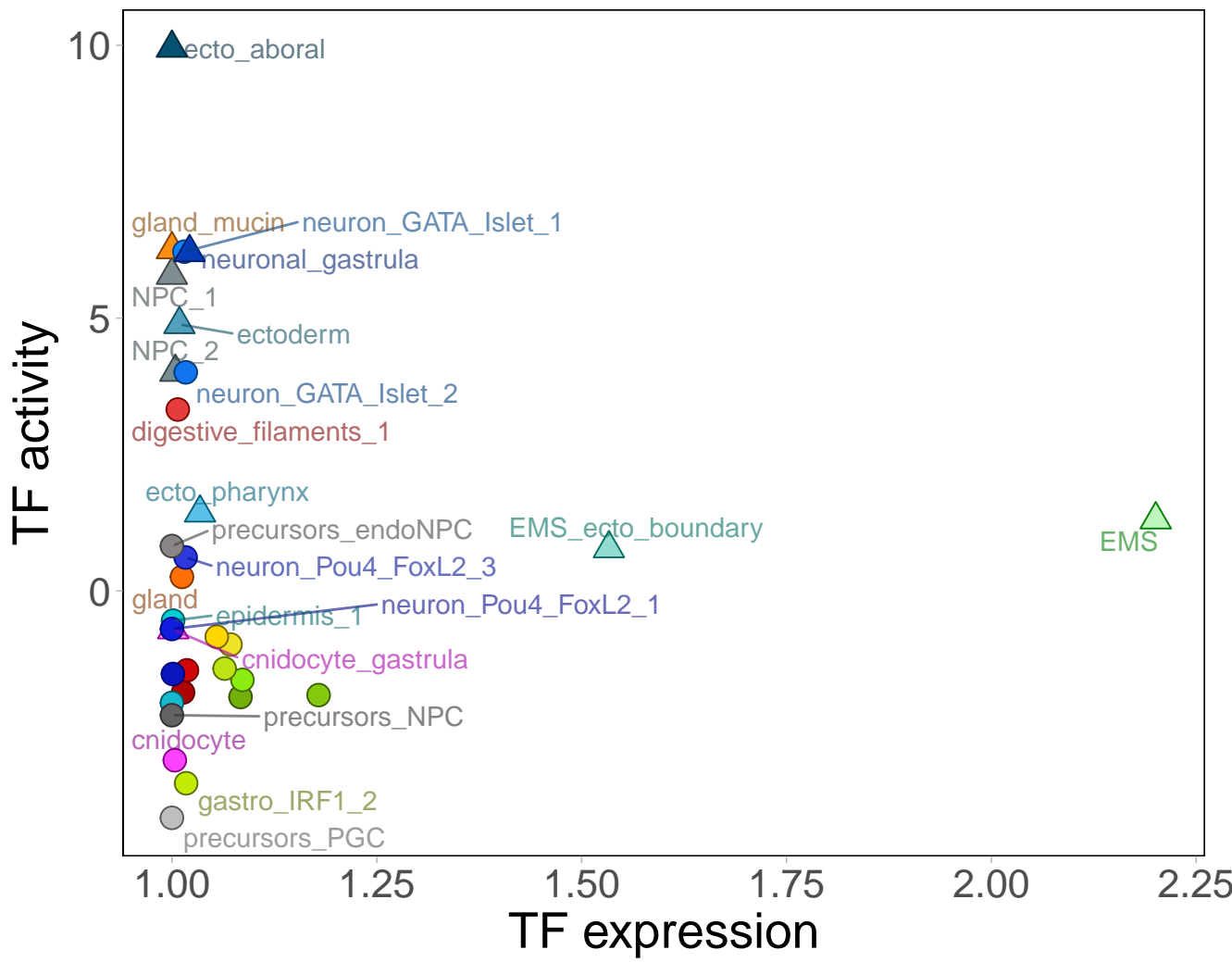


nvec_dap_47
Nvec_vc1.1_XM_032379688.2 (NVHD033-ANTP)
OG: Homeodomains.HG1.39:BARHL1/BARHL2
PFAM: Homeodomain

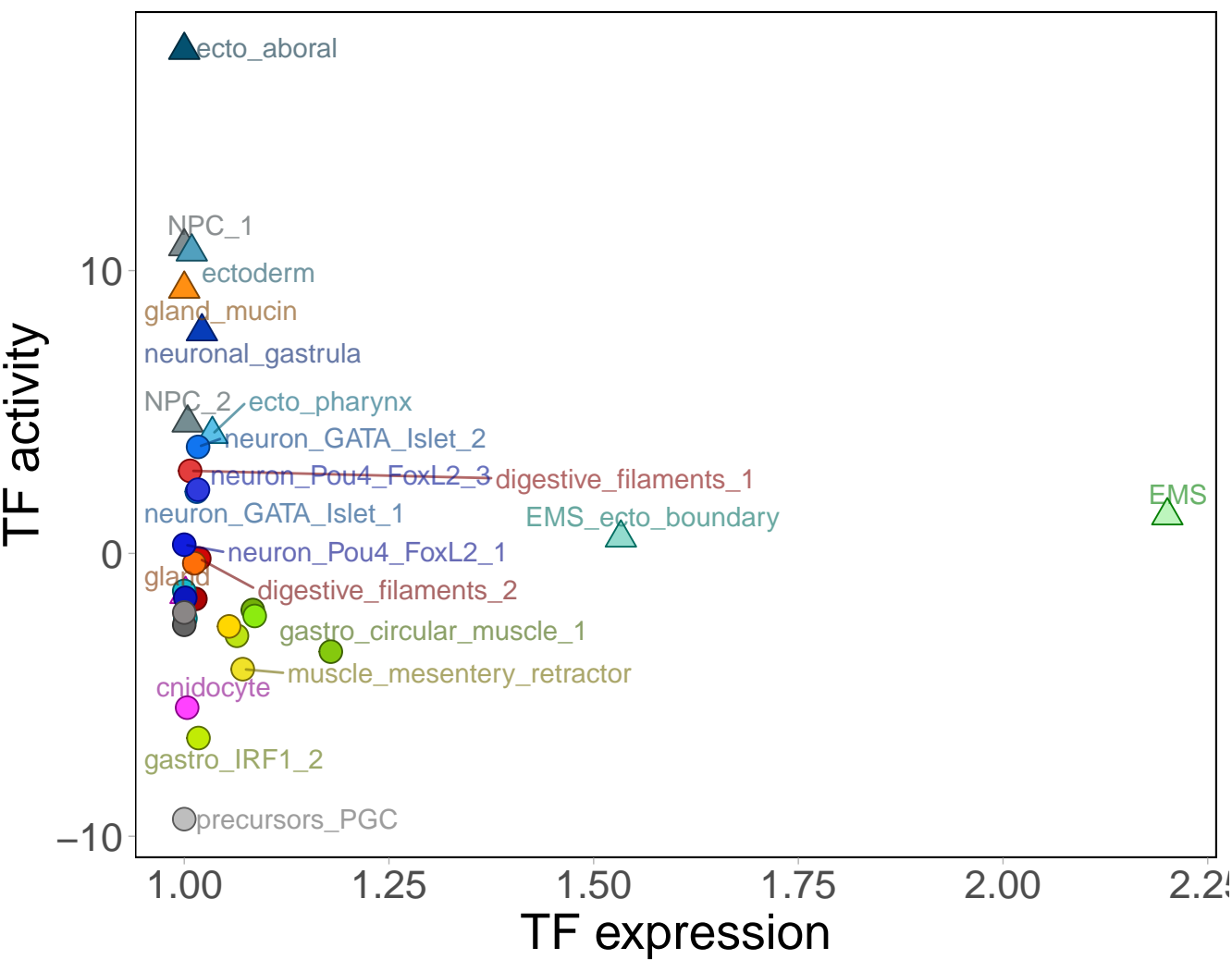
archetype



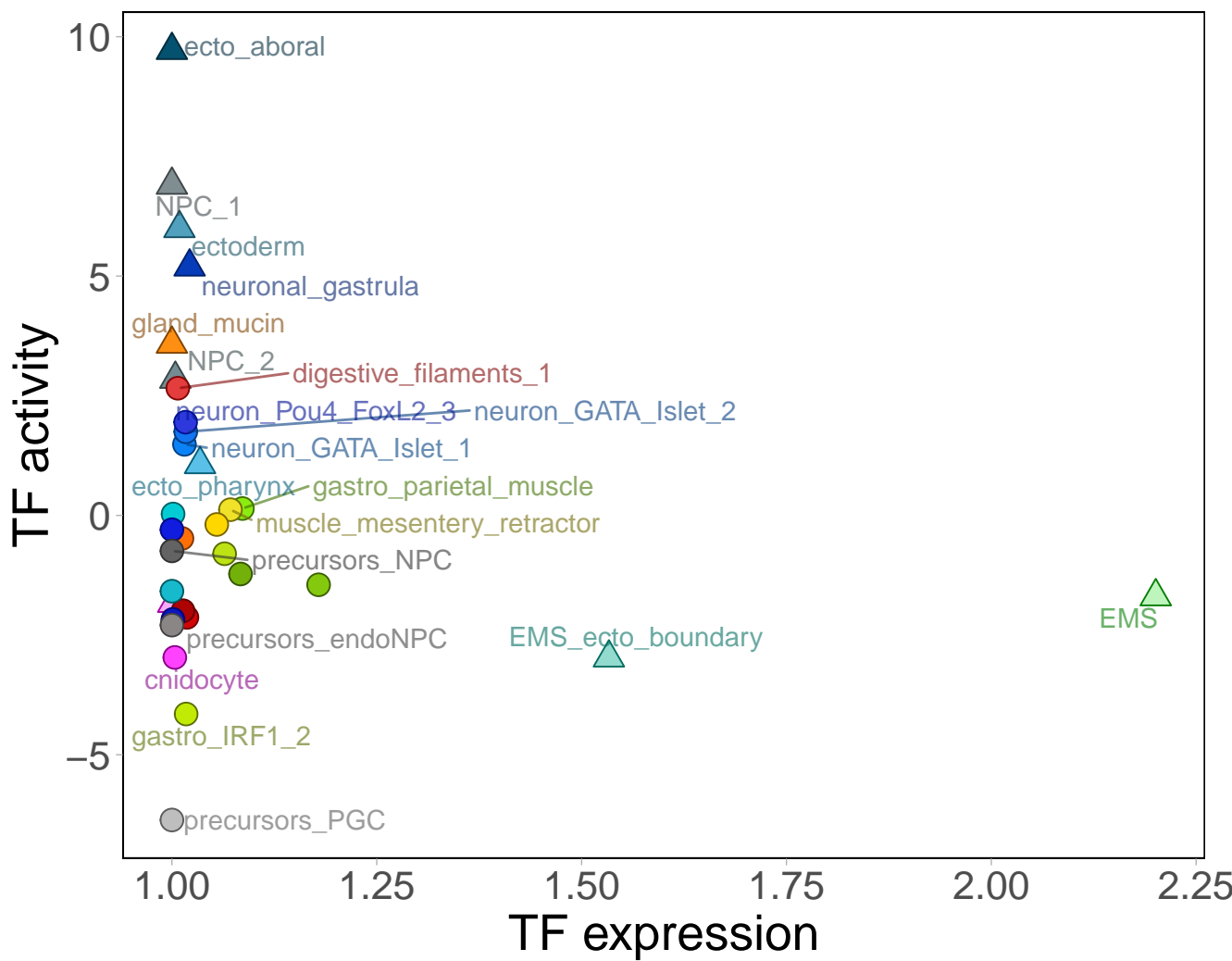
dimont



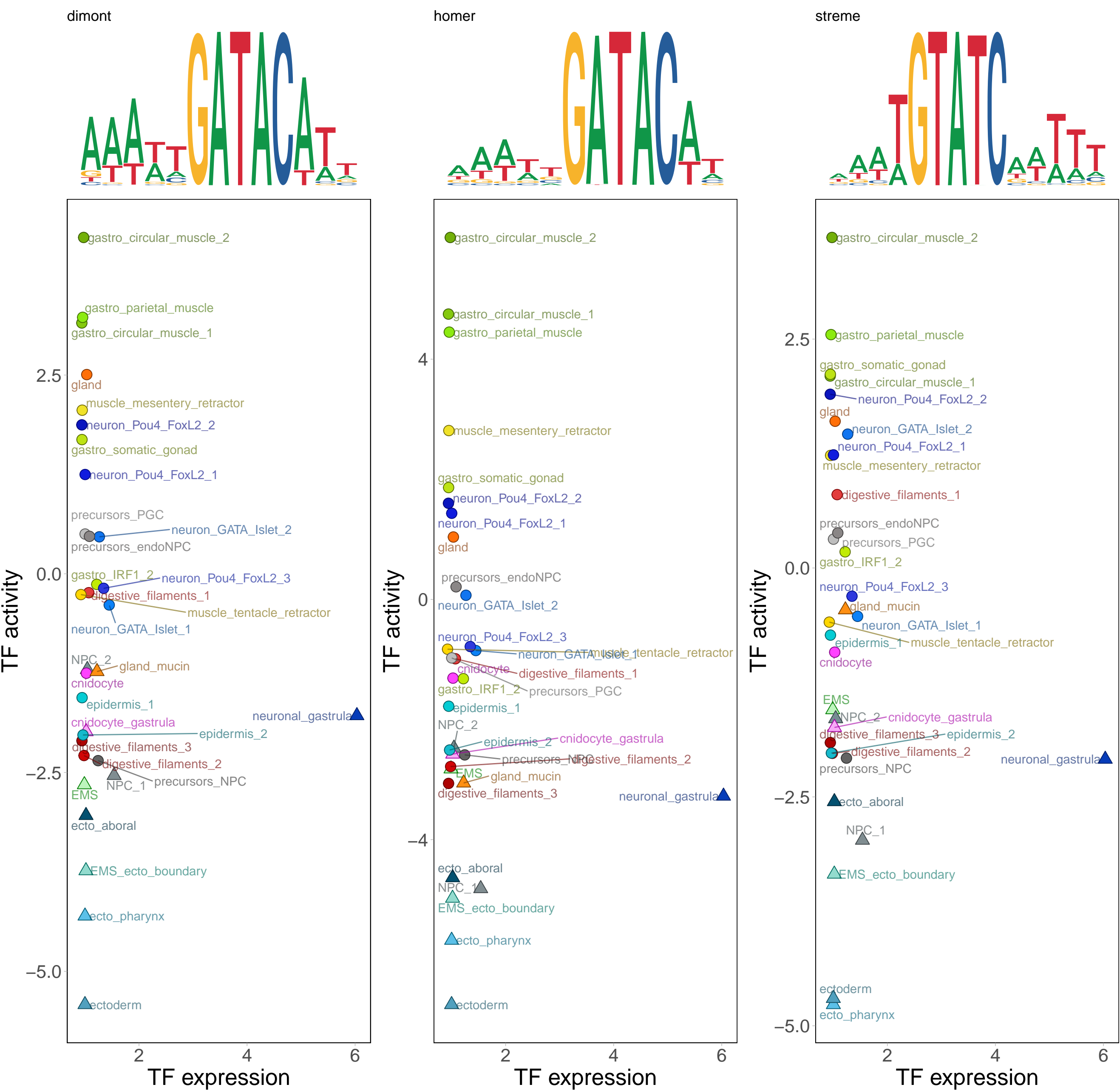
homer



streame

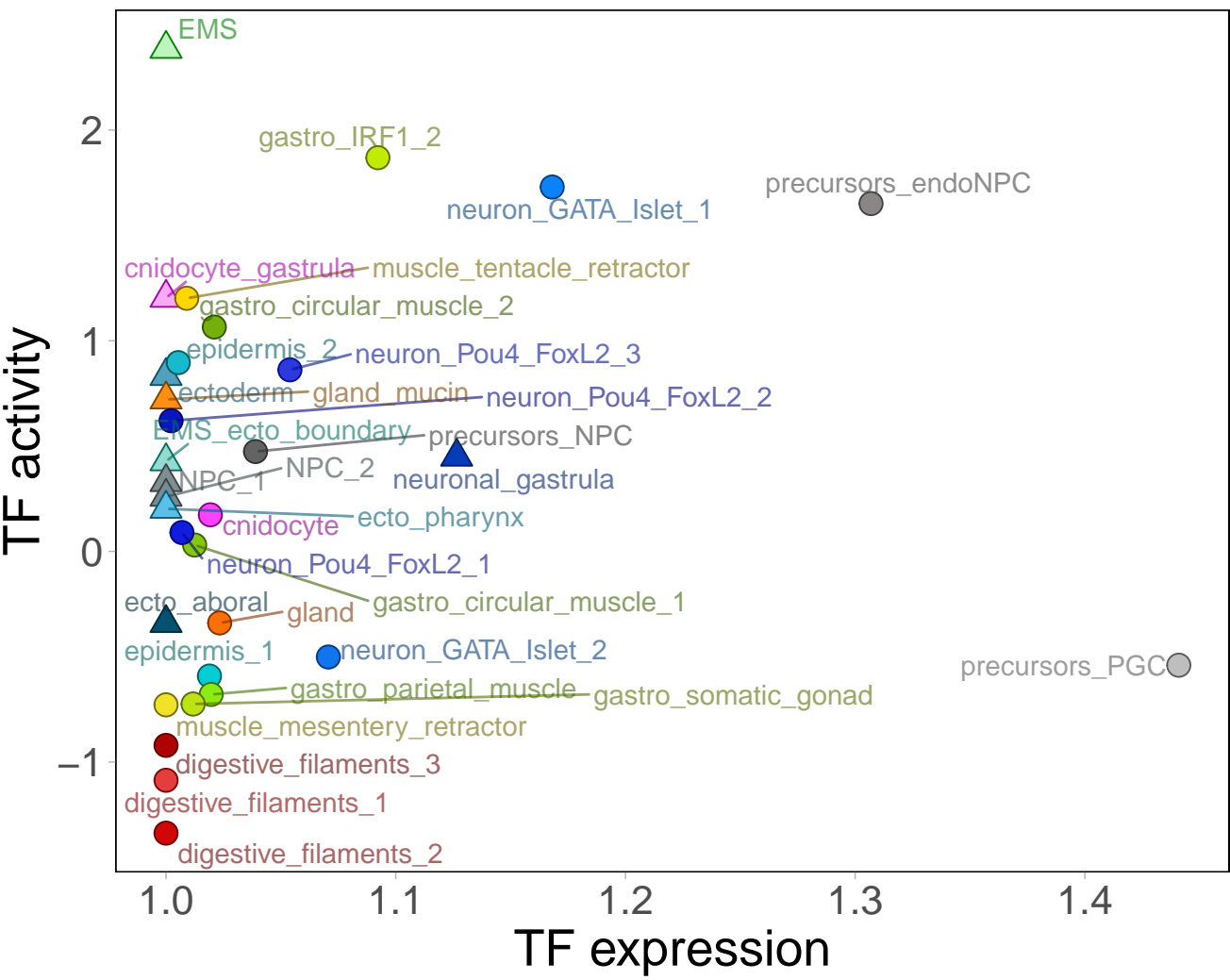


nvec_dap_51
Nvec_vc1.1_XM_001628087.3
OG: Doublesex_DM.HG1.12:DMRTA1/DMRTA2/DMRTB1
PFAM: DM/DMA

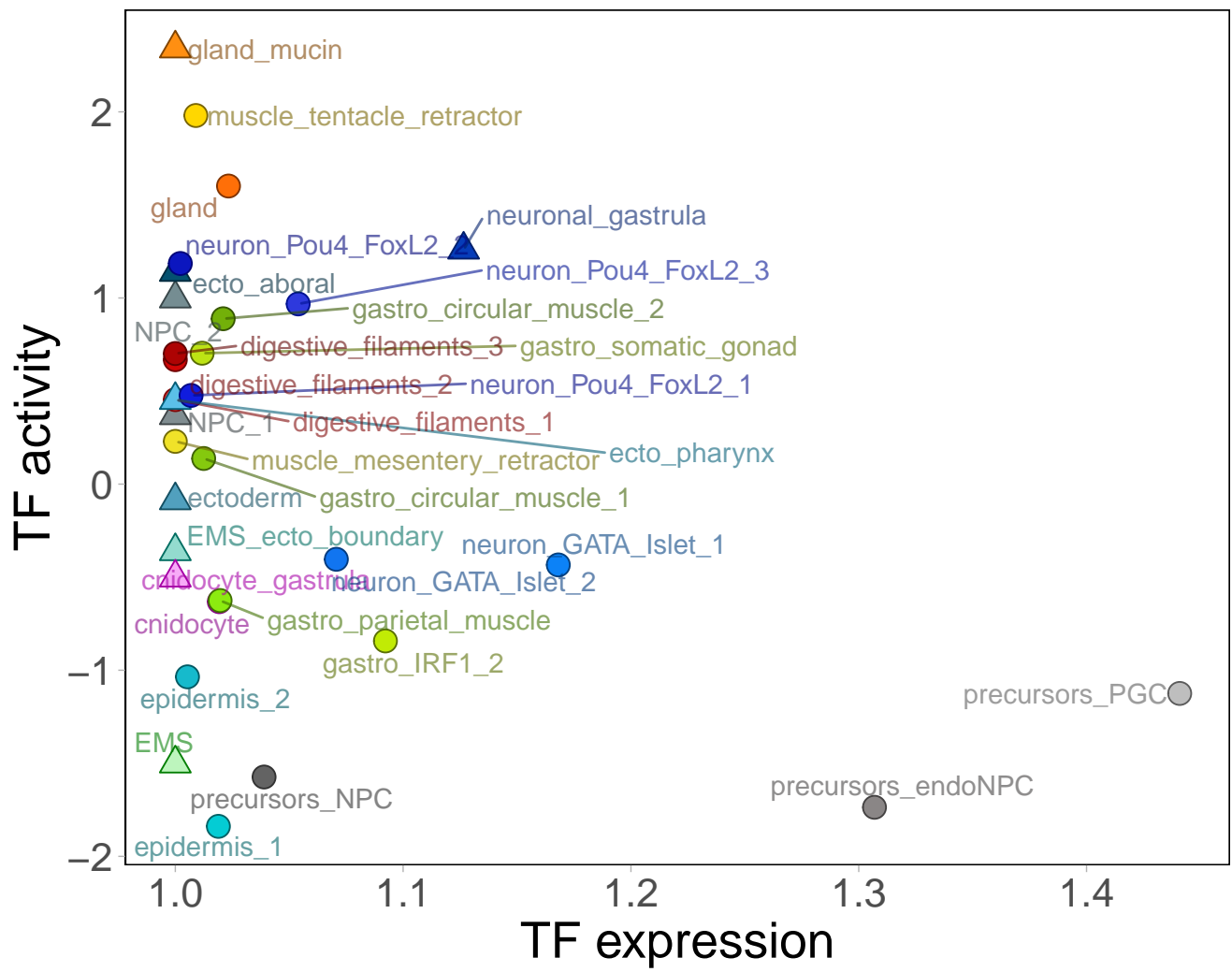


nvec_dap_52
Nvec_vc1.1_XM_032385745.2 (Prdm14d)
OG: zf-C2H2.HG19.1:PRDM14
PFAM: zf-C2H2/zf-C2H2

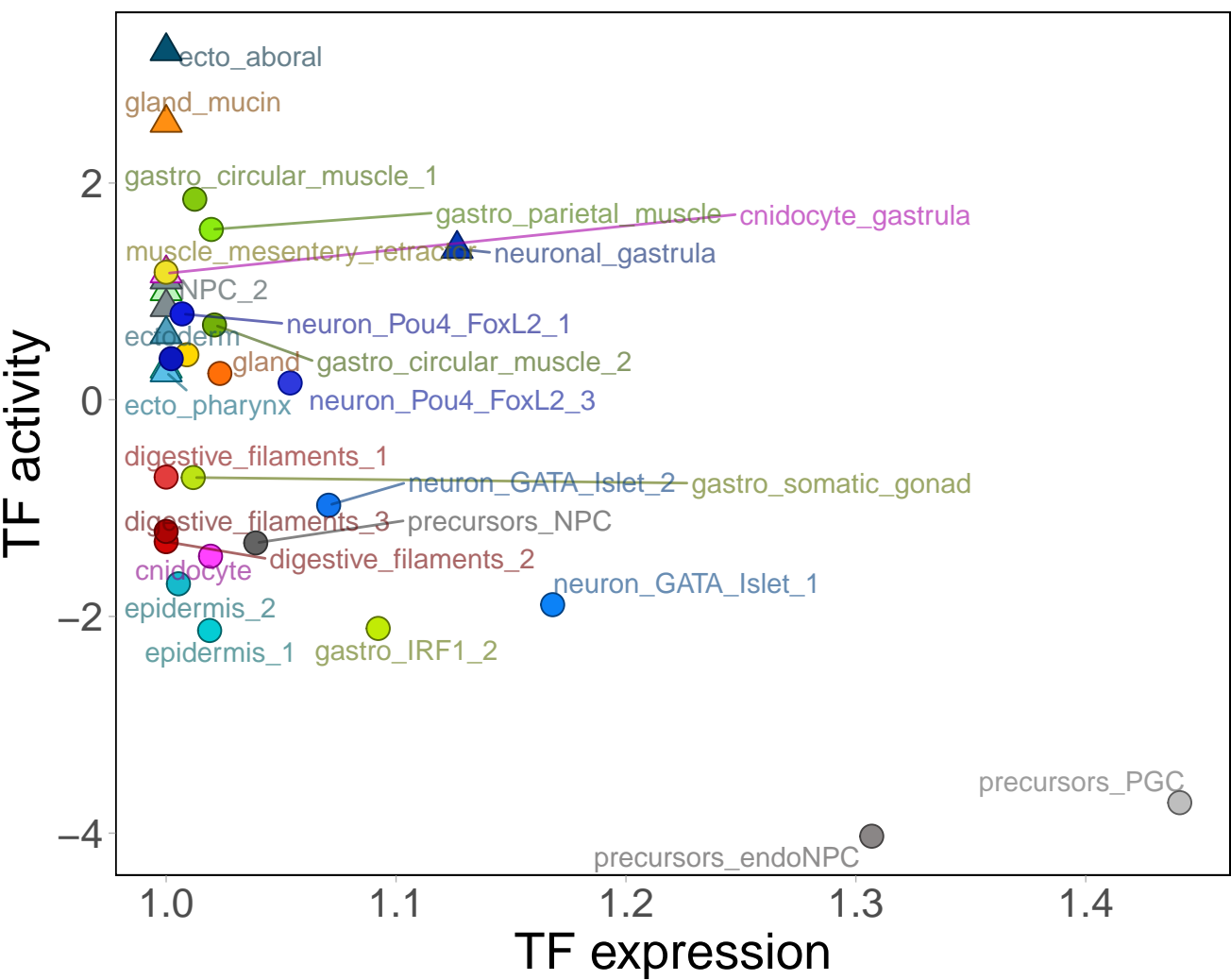
archetype



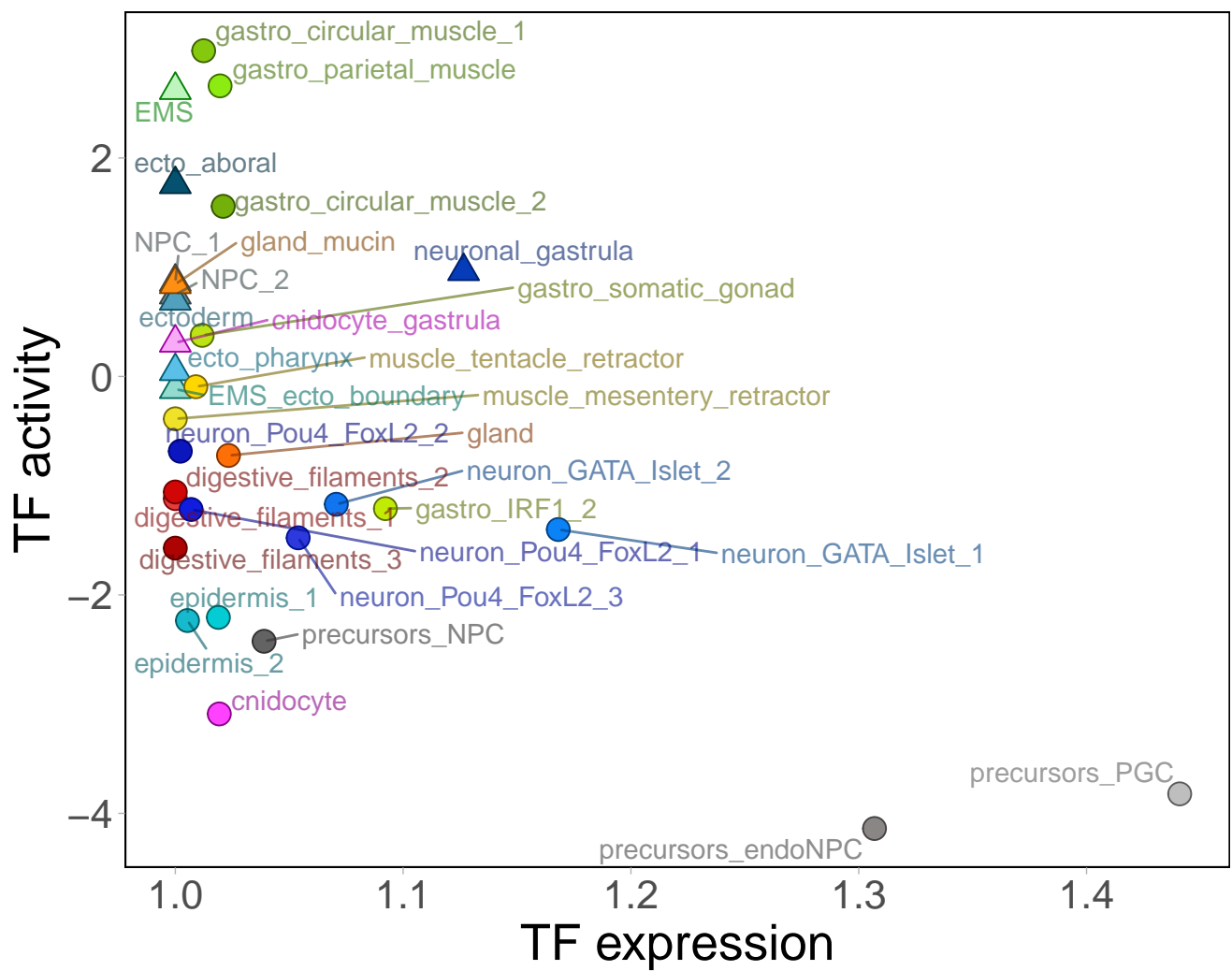
dimont



homer



streame



FoxQ_S5
Nvec_vc1.1_XM_032385706.2 (FoxQ2d)
OG: Forkhead.HG1.14:like:FOXN1/FOXN2/FOXN3/FOXN4/FOXR1/FOXR2
PFAM: Forkhead

