hyp1V_and_ant_arc_V: Hypodermis and seam

C. briggsae TPM
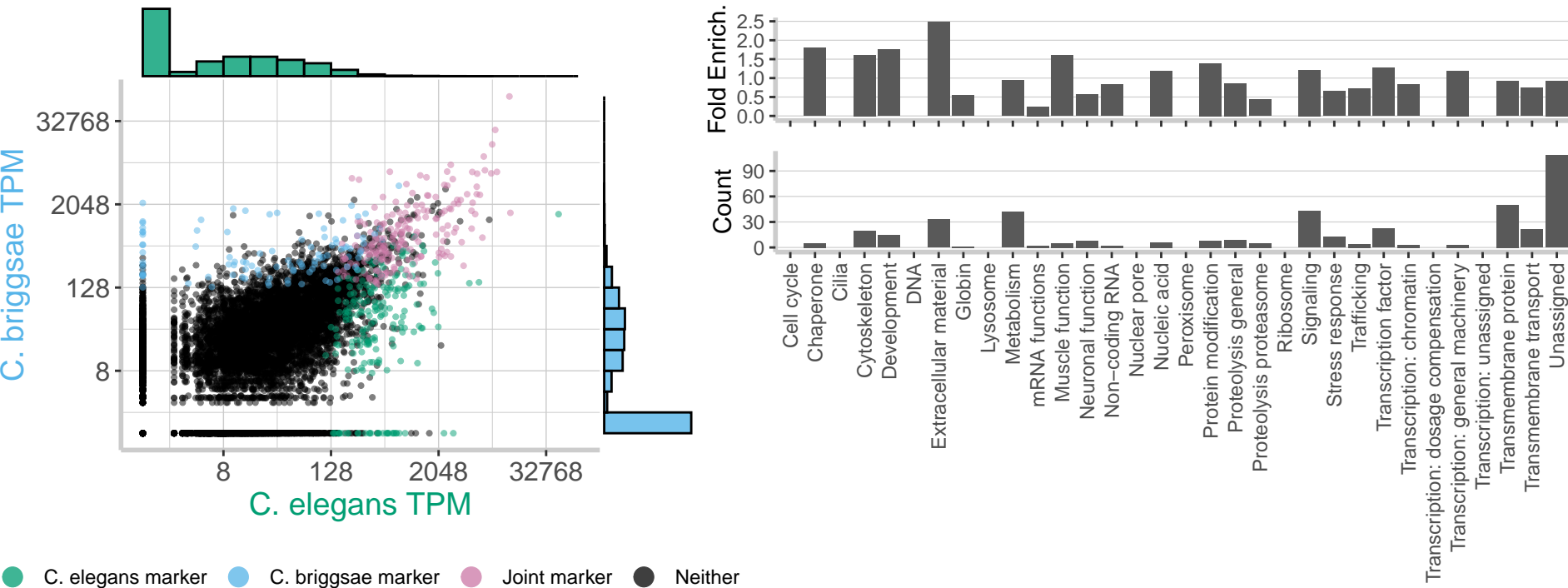
C. elegans TPM

- ● C. elegans marker
- ● C. briggsae marker
- ● Joint marker
- ● Neither

Fold Enrich.

Count

Worm Category (Tier 1)

Cell cycle, Chaperone, Cilia, Cytoskeleton, Development, DNA, Extracellular material, Globin, Lysosome, Metabolism, mRNA functions, Muscle function, Neuronal function, Non-coding RNA, Nuclear pore, Nucleic acid, Peroxisome, Protein modification, Proteolysis general, Proteolysis proteasome, Ribosome, Signaling, Stress response, Trafficking, Transcription factor, Transcription: chromatin, Transcription: dosage compensation, Transcription: general machinery, Transcription: unassigned, Transmembrane protein, Transmembrane transport, Unassigned

TPM: 32768, 4096, 512, 64

C. briggsae / C. elegans

Y54F10AM.6, C30G12.4, Y58A7A.2, cutl-27, C16B8.2, cutl-17, wrt-6, hpo-36, grd-16, qua-1, swip-10, grl-14, col-89, pqn-35, F01G10.9, F56H9.6, T21B4.3, F19H8.2, nlp-48, F33D4.6, CBG08532, K08D12.7, CBG01909, CBG26164, Y6B3B.7

Cell count

Gini Coef.

Genes detected

Ratio of private markers

Mean number of UMI

Jensen–Shannon Distance

Pearson Correlation

Btwn. species DEG