Α

| | | Copepoda | | | | | . | | | | _ | | | |
|-----------|------------------------|----------|---------|----------|-------|------------|-----------|--------|-------|--------|----------|--------|---------|--------|
| Family | Cluster in trees | Calan | Cyclo | Harpa | Sipho | Copep | Crust | Arthr | Ecdys | Lopho | Cepha | Chord | Ambul | Cnidar |
| 2X family | P2X | 44 | 11 | 15 | 6 | 76 | 8 | 3 | 1 | 3 | 1 | 13 | 6 | 4 |
| | ASIC BASIC DEL10 | 1 | | | | 1 | 1 | | 1 | 24 | 10 16 | 9 2 | 10 3 | 21 |
| | DEL10_EGAS1-4 | | ' I | 1 | | 1 | - | | 3 | 1 | 2 | | | |
| | EGAS1-4 | 93 | 29 | 36 | 23 | 181 1 | 153 1 | 01 | 4 | | | | | ć |
| ASIC/Deg/ | ENAC | 35 | 20 | 40 | 11 | 106 | 61 | 32 | 2 | 6 | 14 | 8 | 34 | |
| ENaC | ENAC_ACD_DELM | | | | | | | | 8 | | | | | |
| family | ENAC_degt-1 | | | | | | | | 3 | | | 1 | | |
| | ENAC_ACD_DELM_degt-1 | 45 | 26 | 20 | 3 | 94 | 25 | | 1 | | | | | |
| | FaNaC | 4 | | | | 4 | | | 7 | 25 | | | | |
| | MEC10 | | - | | | | • | | 2 | | | | | |
| | MEC4_MEC10_DEL1 | | | | | | | | 9 | | | | | |
| | PPK | | 281 1 | | | 896 | 15 | 34 | | | | | | |
| | AMPA | 77 | 8 | 17 | 5 | 107 | 7 | 13 | 3 | 23 | 8 | 8 | 2 | |
| | AMPA_GLR1-2 | | | | | | | | 2 | | | | | |
| | AMPA_GluRIA | 16 | 4 | 5 | 6 | 31 | 5 | 13 | | | | | | |
| | Delta | | | | | | | | | | 29 | 4 | 3 | |
| | Epsilon | | | | | | | | | | 7 | | | ! |
| | KAINATE | 201 | 35 | 41 | 58 | 335 | 24 | 58 | 5 | 7 | 2 | 10 | 1 | |
| | KAINATE_GLR-3 | | | | | | | | 2 | | | | | |
| | NMDA1 | 13 | 6 | 3 | 6 | 28 | 1 | 1 | 2 | 3 | 2 | 2 | 2 | |
| | NMDA2 | 32 | 12 | 9 | 16 | 69 | 9 | 8 | 3 | 7 | 5 | 7 | | |
| | NMDA3 | | | | | | 2 | 3 | 1 | 7 | 2 | 4 | 2 | |
| | iGluR* | 2 | | | 10 | 2 | | | , 1 | 20 | 12 | | | |
| | IR8 | 35 | 16 | 22 | 18 | 91 | 3 | 3 | | | | | | |
| iGluR | IR25 | 43 | 8 | 7 | 8 | 66 | 4 | 9 | 2 | 2 3 | | | | |
| family | IR25a-like_8a-like | 20 | 7 | | 20 | F 7 | 200 | 2.6 | 1.0 | | | | | |
| | IR* | 28 | 7 | 2 | 20 | | 298 | 26 | 18 | 28 | | | | |
| | IR21a | 6 | | 12 | 7 | 13 | 1 | 2 | | | | | | |
| | IR321_IR335 | 23 | 1 | 13 | 16 | 53 | 2 | 1 | | | | | | |
| | IR324_IR68a | 27 | 3 | 5 | 9 | 44 | 21 | 1 | | | | | | |
| | IR334 | 68 80 | 6 30 | 32 55 | | 113 190 | 4 36 | 10 | ı | | | | | |
| | IR336 | 80 | | | | | 36 | 48 | | | | | | |
| | IR337 | | 6 | 5 | 16 | 27 | _ 0_ | | 1 | | | | | |
| | IR40a | | _ 1_ | | | _ 1_ | 8 | 3 | | | | | | |
| | IR75a | . 1 | 1 | | | 1 | | 7 | | | | | | |
| | IR76a_IR41a | 1 | | | | 1 | | 4 | | | | | | |
| | IR76b | 9 | 2 | 3 | 1.0- | 14 | 9 | 2 6 | | | | | | |
| | IR93a | 40 | 3 | 6 | 10 | 59 | 0 | 6 | | | | | | |

B

| Cepha Chord Ambul Cnidar | Family | Cluster in trees | Cyclo Cyclo Bpodadoo | Copep | Crust | Ecdys | Lopho | Chord | Ambul |
|--|---|--|---|------------------------------|--------------------------|---|---|----------------------------------|-----------------|
| 1 13 6 4 10 9 10 21 16 2 3 2 9 14 8 34 7 1 1 8 8 2 29 4 3 9 2 10 1 2 2 2 7 | Cys-loop family: Acetylcholine and 5HT receptor | 5HTR_A-E 5HTR-like AChR* AChR_alpha_1-6 AChR_alpha_1-like AChR_alpha_2-like AChR_alpha_7 AChR_alpha_9-10 AChR_beta_1-like AChR_beta_2-like AChR_beta_2-like Anion-selective LnAChR B-I-F-K Cation-selective LnAChR D Cation-selective LnAChR D Cation-selective LnAChR G Cation-selective LnAChR G Cation-selective LnAChR H Cation-selective LnAChR J des-2_acr-23_deg-3_acr-5 PBO_5-6 unc-29_lev-1_acr-2-3 | 19 4 6 9 10 3 7 6 | 26 194 36 | 4 6 | 30 8 5 6 12 13 6 7 6 3 13 19 3 1 | 18 5 11 2 3 9 10 3 | 7 7 15 7 15 1 4 39 4 | 72 50 2 1 |
| 5 7 3 2 4 2 12 | Cys-loop family: GABA and Glycine channel | Anionic_ACC/Monoamine- gated_channel Anionic_glutamate Cys_loop* GABA GABA_alpha_gama GABA_beta_theta GABA_EXP_1 GABA_pi_delta GABA_RDL GABA_rho GABA_UNC-49B GlyR-like GlyR_alpha GlyR_beta Histamine-gated chloride channel_alpha_1 lgc-57_GRR pH sensitive chloride channel 2 pH sensitive chloride channel 1 | 14 6 2 1 99 23 17 23 18 5 6 8 97 30 29 9 | 165 10 349 72 15 | 3 12 5 11 40 | 16 23 8 1 11 4 6 3 2 19 1 2 38 15 11 7 9 6 | 11 5 11 22 1 | 7 | 42 1 43 |