

| GO Term | Description | FDR q-value |
|------------|--|-------------|
| GO:0044421 | extracellular region part | 7.73E-12 |
| GO:0005576 | extracellular region | 1.44E-11 |
| GO:0005581 | collagen trimer | 8.85E-11 |
| GO:0005615 | extracellular space | 1.03E-09 |
| GO:0031012 | extracellular matrix | 2.58E-07 |
| GO:0032502 | developmental process | 3.22E-07 |
| GO:0044767 | single-organism developmental process | 6.04E-07 |
| GO:0048646 | anatomical structure formation involved in morphogenesis | 6.57E-07 |
| GO:0044707 | single-multicellular organism process | 8.78E-06 |
| GO:0032501 | multicellular organismal process | 1.10E-05 |
| GO:0098797 | plasma membrane protein complex | 1.90E-05 |
| GO:0044459 | plasma membrane part | 2.25E-05 |
| GO:0044724 | single-organism carbohydrate catabolic process | 2.48E-05 |
| GO:0007155 | cell adhesion | 2.60E-05 |
| GO:0016021 | integral component of membrane | 2.69E-05 |
| GO:0044699 | single-organism process | 2.87E-05 |
| GO:0022610 | biological adhesion | 2.97E-05 |
| GO:0007166 | cell surface receptor signaling pathway | 3.75E-05 |
| GO:0048856 | anatomical structure development | 4.04E-05 |
| GO:0004872 | receptor activity | 4.57E-05 |
| GO:0031224 | intrinsic component of membrane | 4.68E-05 |
| GO:0016052 | carbohydrate catabolic process | 6.86E-05 |
| GO:0001525 | angiogenesis | 7.27E-05 |