

<b>Annotation Cluster 1</b>		Enrichment Score: 1.89	
<b>GO term</b>	<b>Count</b>	<b>P_Value</b>	
cell-cell adhesion	7	3.60E-03	
cell adhesion	9	2.40E-02	
biological adhesion	9	2.40E-02	

<b>Annotation Cluster 2</b>		Enrichment Score: 1.81	
<b>GO term</b>	<b>Count</b>	<b>P_Value</b>	
regulation of neurogenesis	5	9.40E-03	
regulation of nervous system development	5	1.40E-02	
regulation of cell development	5	1.80E-02	
regulation of neuron differentiation	4	2.60E-02	

<b>Annotation Cluster 1</b>		Enrichment Score: 4.11	
<b>GO term</b>	<b>Count</b>	<b>P_Value</b>	
tube morphogenesis	10	3.80E-05	
tube development	12	4.60E-05	
epithelium development	11	2.70E-04	

<b>Annotation Cluster 2</b>		Enrichment Score: 3.63	
<b>GO term</b>	<b>Count</b>	<b>P_Value</b>	
epithelial tube morphogenesis	8	9.30E-05	
morphogenesis of an epithelium	9	2.60E-04	
epithelium development	11	2.70E-04	
tissue morphogenesis	10	4.70E-04	

<b>Annotation Cluster 3</b>		Enrichment Score: 3.33	
<b>GO term</b>	<b>Count</b>	<b>P_Value</b>	
chordate embryonic development	14	2.00E-04	
embryonic development ending in birth or egg hatching	14	2.20E-04	
embryonic morphogenesis	11	2.40E-03	

<b>Annotation Cluster 4</b>		Enrichment Score: 2.51	
<b>GO term</b>	<b>Count</b>	<b>P_Value</b>	
primary neural tube formation	5	6.80E-04	
neural tube formation	5	1.40E-03	
embryonic epithelial tube formation	5	1.70E-03	
tube lumen formation	5	2.40E-03	
neural tube closure	4	6.10E-03	
tube closure	4	6.10E-03	
morphogenesis of embryonic epithelium	5	6.60E-03	
neural tube development	5	8.20E-03	

<b>Annotation Cluster 5</b>		Enrichment Score: 1.96	
<b>GO term</b>	<b>Count</b>	<b>P_Value</b>	
embryonic skeletal system morphogenesis	5	3.30E-03	
skeletal system morphogenesis	6	8.20E-03	
embryonic skeletal system development	5	8.20E-03	
embryonic organ morphogenesis	5	6.90E-02	

<b>Annotation Cluster 6</b>		Enrichment Score: 1.84	
<b>GO term</b>	<b>Count</b>	<b>P_Value</b>	
cell morphogenesis	9	9.70E-03	
cell morphogenesis involved in differentiation	7	1.60E-02	
cellular component morphogenesis	9	2.00E-02	

<b>Annotation Cluster 7</b>		Enrichment Score: 1.52	
<b>GO term</b>	<b>Count</b>	<b>P_Value</b>	
pyrimidine nucleotide biosynthetic process	4	8.70E-04	
pyrimidine nucleotide metabolic process	4	2.10E-03	
nucleotide biosynthetic process	5	8.60E-02	
nucleobase, nucleoside and nucleotide biosynthetic process	5	9.30E-02	
nucleobase, nucleoside, nucleotide and nucleic acid biosynthetic process	5	9.30E-02	
nucleoside metabolic process	3	1.00E-01	
nitrogen compound biosynthetic process	6	1.60E-01	