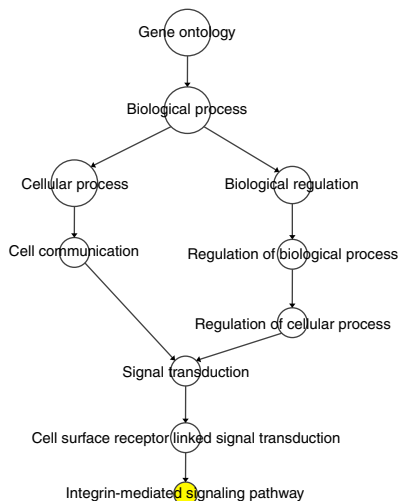
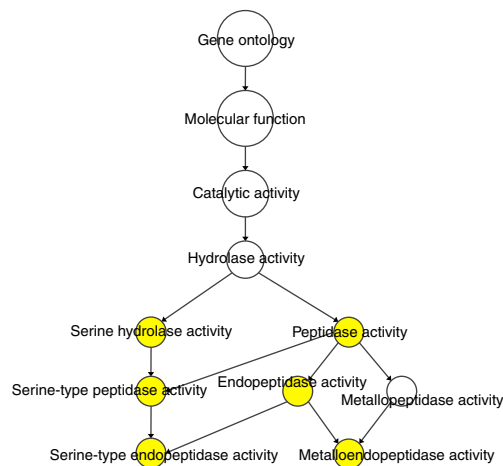


(a)**(b)****(c)**

Term ID	Term	Original <i>P</i> -value	Corrected <i>P</i> -value
Biological process:			
GO:7229	Integrin-mediated signaling pathway	3.63×10^{-05}	1.03×10^{-02}
Molecular function:			
GO:0016809	Endopeptidase activity	2.97×10^{-05}	5.94×10^{-03}
GO:0004222	Metalloendopeptidase activity	6.74×10^{-04}	4.14×10^{-02}
GO:0008233	Peptidase activity	7.61×10^{-04}	4.14×10^{-02}
GO:0004252	Serine-type endopeptidase activity	8.27×10^{-04}	4.14×10^{-02}
GO:0008236	Serine-type peptidase activity	1.45×10^{-03}	4.93×10^{-02}
GO:0017171	Serine hydrolase activity	1.48×10^{-03}	4.93×10^{-02}
Pfam:			
PF01562	Reprolysin family propeptide	1.78×10^{-05}	4.24×10^{-04}
PF01421	Reprolysin (M12B) family zinc metalloprotease	2.15×10^{-05}	4.24×10^{-04}