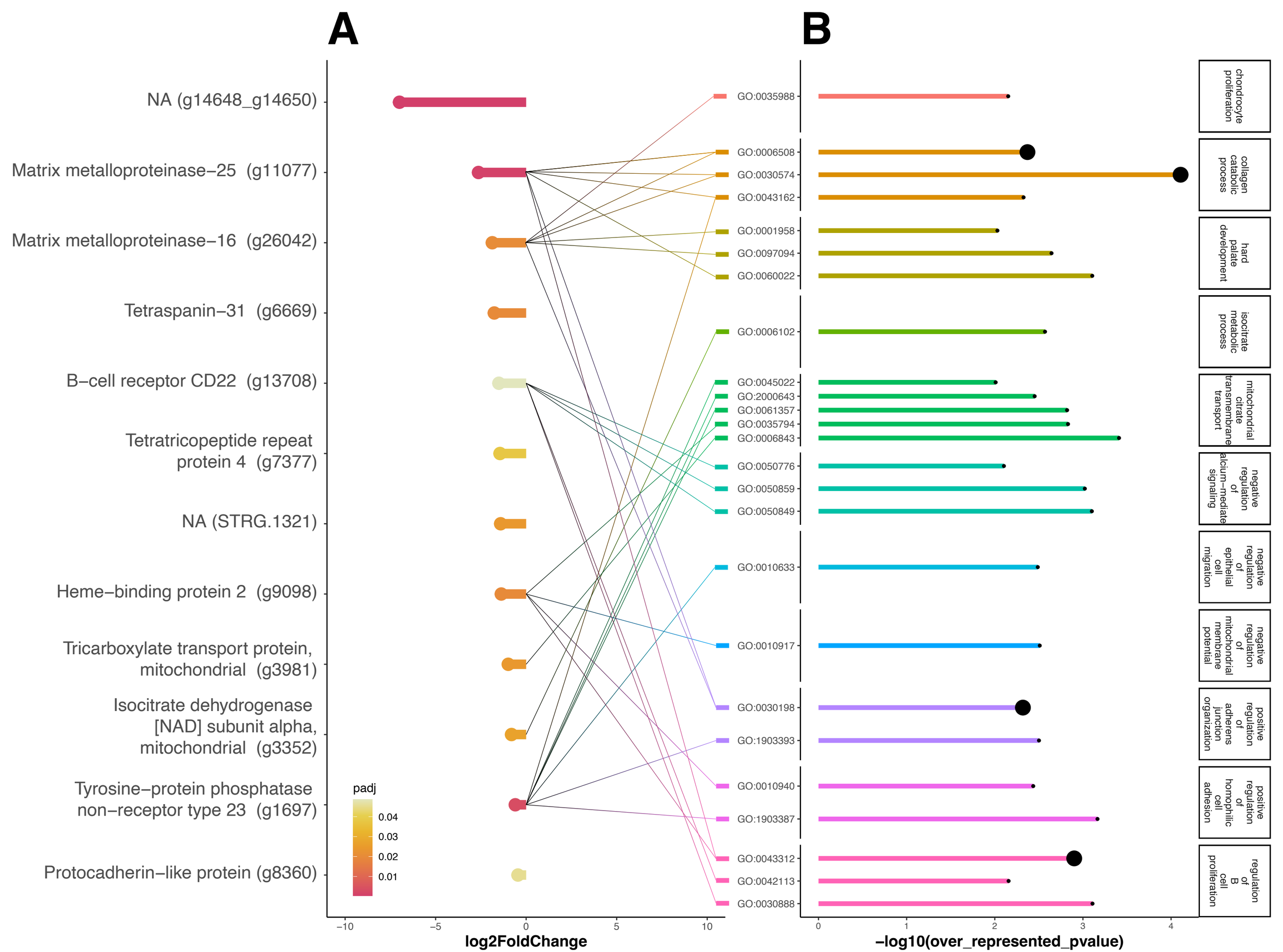
A
B
NA (g14648_g14650)
Matrix metalloproteinase−25 (g11077)
Matrix metalloproteinase−16 (g26042)
Tetraspanin−31 (g6669)
B−cell receptor CD22 (g13708)
Tetratricopeptide repeat protein 4 (g7377)
NA (STRG.1321)
Heme−binding protein 2 (g9098)
Tricarboxylate transport protein, mitochondrial (g3981)
Isocitrate dehydrogenase [NAD] subunit alpha, mitochondrial (g3352)
Tyrosine−protein phosphatase non−receptor type 23 (g1697)
Protocadherin−like protein (g8360)
padj
0.04
0.03
0.02
0.01
−10
−5
0
5
10
log2FoldChange
GO:0035988
GO:0006508
GO:0030574
GO:0043162
GO:0001958
GO:0097094
GO:0060022
GO:0006102
GO:0045022
GO:2000643
GO:0061357
GO:0035794
GO:0006843
GO:0050776
GO:0050859
GO:0050849
GO:0010633
GO:0010917
GO:0030198
GO:1903393
GO:0010940
GO:1903387
GO:0043312
GO:0042113
GO:0030888
chondrocyte proliferation
collagen catabolic process
hard palate development
isocitrate metabolic process
mitochondrial citrate transmembrane transport
negative regulation of calcium−mediated signaling
negative regulation of epithelial cell migration
negative regulation of mitochondrial membrane potential
positive regulation of adherens junction organization
positive regulation of homophilic cell adhesion
regulation of B cell proliferation
0
1
2
3
4
−log10(over_represented_pvalue)