

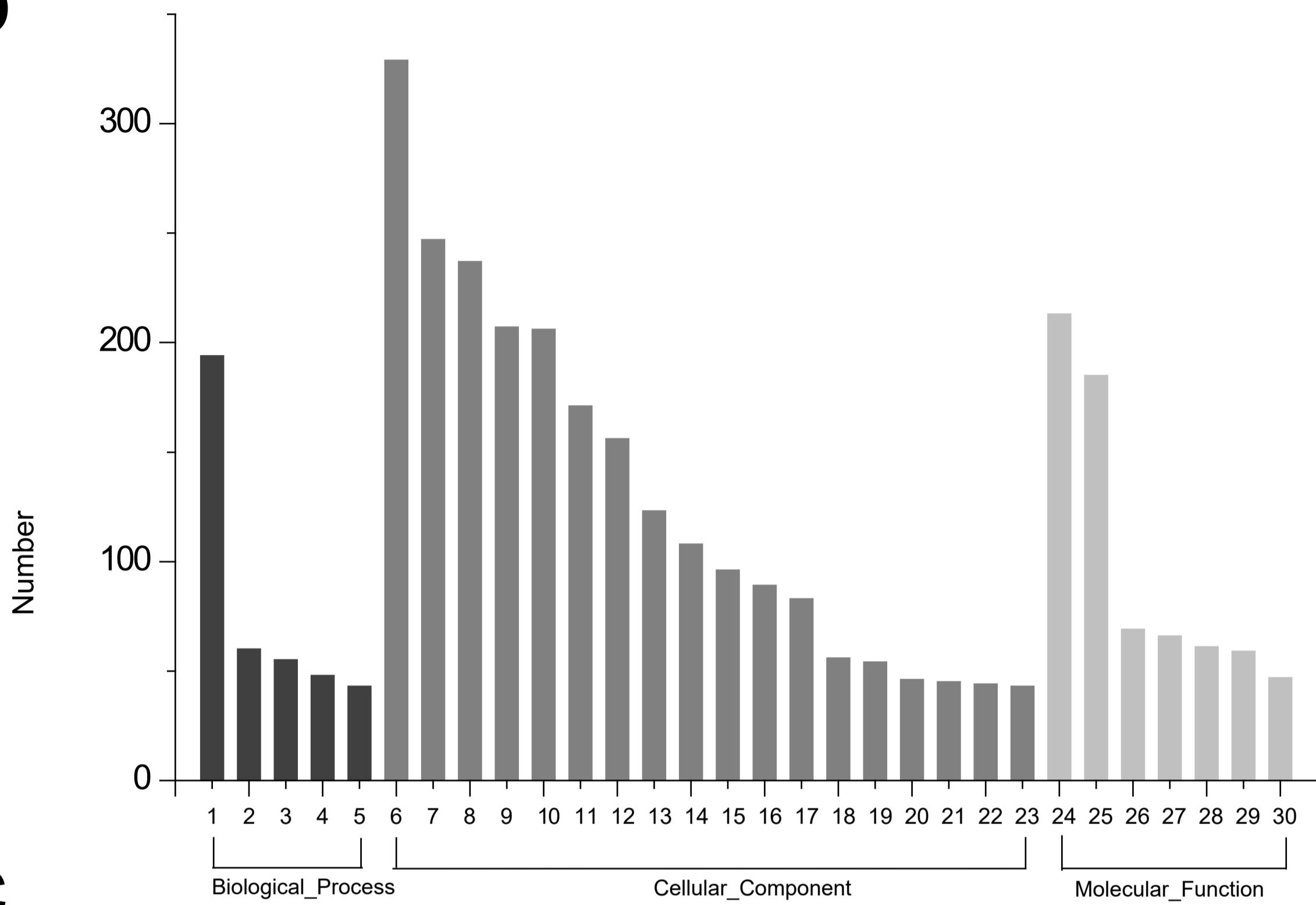
a

L1_profile23 L2_profile23

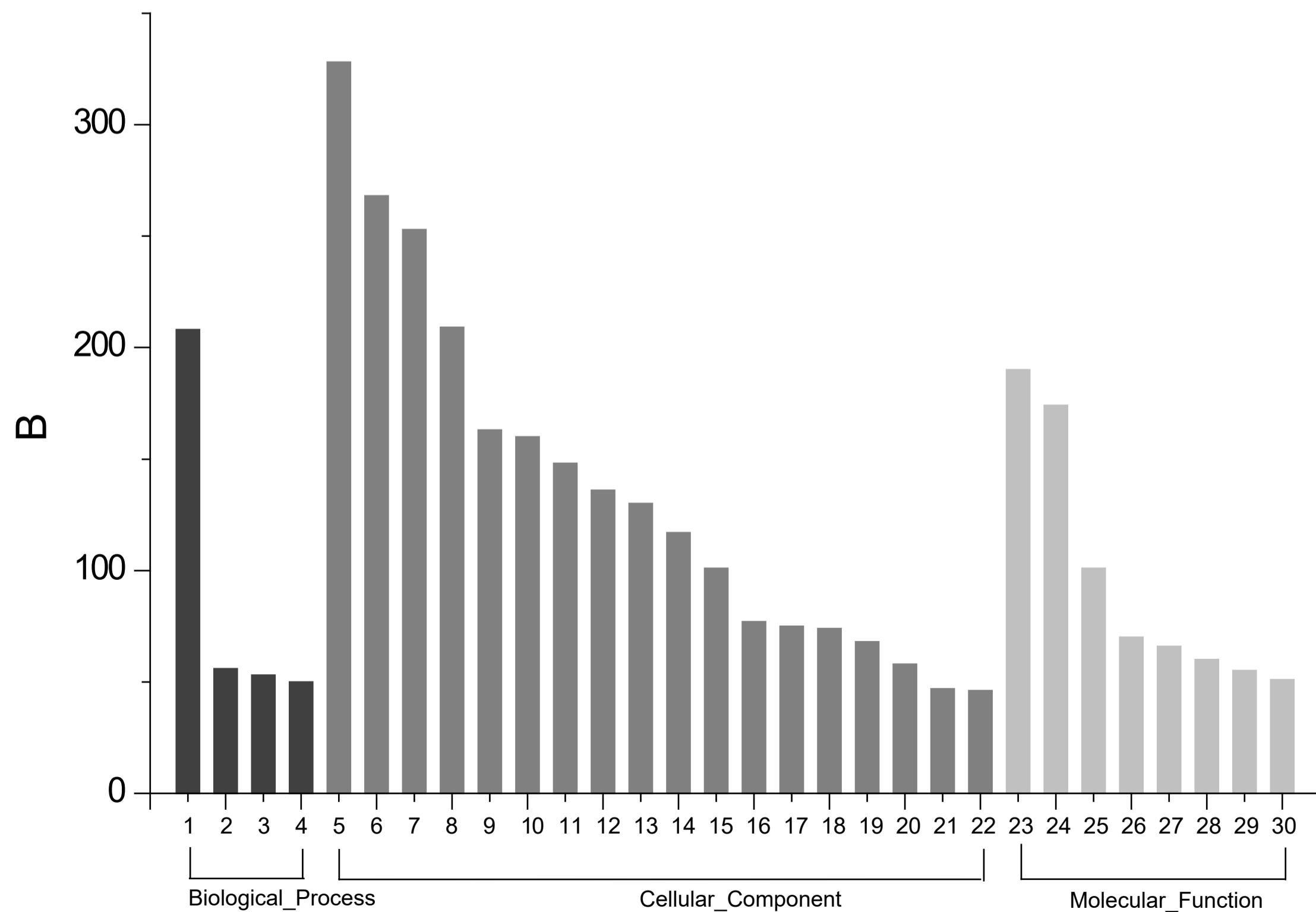
723

189

689

b

- 1:biological_process
- 2:response to salt stress
- 3:regulation of transcription, DNA-templated
- 4:transcription, DNA-templated
- 5:oxidation-reduction process
- 6:nucleus
- 7:cytoplasm
- 8:plasma membrane
- 9:chloroplast
- 10:integral component of membrane
- 11:cytosol
- 12:membrane
- 13:mitochondrion
- 14:Golgi apparatus
- 15:extracellular region
- 16:plasmodesma
- 17:cellular_component
- 18:vacuole
- 19:vacuolar membrane
- 20:endosome
- 21:endoplasmic reticulum
- 22:Golgi membrane
- 23:cell wall
- 24:protein binding
- 25:molecular_function
- 26:ATP binding
- 27:metal ion binding
- 28:zinc ion binding
- 29:DNA binding
- 30:transcription factor activity, sequence-specific DNA binding

C

- 1:biological_process
- 2:protein phosphorylation
- 3:regulation of transcription, DNA-templated
- 4:phosphorylation
- 5:nucleus
- 6:plasma membrane
- 7:integral component of membrane
- 8:cytoplasm
- 9:chloroplast
- 10:membrane
- 11:cytosol
- 12:Golgi apparatus
- 13:mitochondrion
- 14:extracellular region
- 15:plasmodesma
- 16:cellular_component
- 17:endosome
- 18:trans-Golgi network
- 19:vacuole
- 20:Golgi membrane
- 21:cell wall
- 22:endoplasmic reticulum
- 23:molecular_function
- 24:protein binding
- 25:ATP binding
- 26:metal ion binding
- 27:kinase activity
- 28:transferase activity, transferring glycosyl groups
- 29:zinc ion binding
- 30:protein serine/threonine kinase activity