|  | interleukin–6 production  positive regulation of interleukin–13 production  interleukin–5 production  positive regulation of interleukin–6 production  regulation of interleukin–6 production  positive regulation of interleukin–5 production  regulation of interleukin–5 production      |   |
|--|---|---|
| response to interleukin–7  response to interleukin–17  response to interleukin–15  Cellular response to interleukin–15  cellular response to interleukin–7 | regulation of amyloid fibril formationo   | number of genes  O 1.00  O 1.25  O 1.50  O 1.75 |
| interleukin–4–mediated signaling pathway interleukin–17–mediated signaling pathway interleukin–15–mediated signaling pathway                               | negative regulation of amyloid fibril formation   | p.adjust 0.04951817                             |
|  | negative regulation of glycolytic process   |   |
| fibroblast activation  | purine ribonucleoside bisphosphate metabolic process cellular biogenic amine catabolic process  amine catabolic process 3'-phosphoadenosine 5'-phosphosulfate metabolic process  primary alcohol catabolic process  phenol-containing compound catabolic process  ethanol catabolic process |   |