
6238 Com Data If Driver ~UPD~

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the genes related to cl in humans are sometimes related to the similar phenotype in mice. therefore, the mouse model with cl may be useful for the study of human cl. genes associated with cl in humans were identified by the systematic review and mgi database search. the identified genes may be used to help develop new therapeutic strategies to treat cl in humans. we identified genes related to cl in mice using a systematic literature review and mgi database search. the systematic review and mgi database search identified a total of 55 genes (table 1). among them, 39 genes were identified in mice with cl/p resulting from a single gene deficiency (table 1 and table 2). there are nine spontaneous cl/p mouse lines (four genes after excluding any duplicated genes; five mouse lines with spontaneous mutations in cl-associated genes and four mouse lines with spontaneous mutations in unknown gene and loci). the penetrance of cl/p in spontaneous mouse lines is quite low (less than 40%) (table 2). ten compound mutant mice (mice with two mutant genes; 12 genes after excluding any duplicated genes) exhibited cl (table 3). among these 55 cl-associated genes, 20.0% (11 out of 55 genes) were common in the systematic review and mgi database search. there were 14 genes (25.5%, 14 out of 55 genes) and 30 genes (54.5%, 30 out of 55 genes) uniquely identified through the systematic review and mgi search, respectively (fig. 2). the genetic factors related to cl in humans are sometimes related to the similar phenotype in mice. therefore, the mouse model with cl may be useful for the study of human cl. genes related to cl in humans were identified by the systematic review and mgi database search. the identified genes may be used to help develop new therapeutic strategies to treat cl in humans. 5ec8ef588b

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