

Table 2. Quantitative analysis of *Mrg* receptors coexpressed within rat DRG neuronsPercentage of cells that coexpress the indicated *Mrgs* relative to the reference gene
± SEM

<i>Mrg</i> *	$\frac{\text{MrgB4+Mrg}^*}{\text{MrgB4}}$	$\frac{\text{MrgB4+Mrg}^*}{\text{Mrg}^*}$	$\frac{\text{MrgC+Mrg}^*}{\text{MrgC}}$	$\frac{\text{MrgC+Mrg}^*}{\text{Mrg}^*}$	$\frac{\text{MrgD+Mrg}^*}{\text{MrgD}}$	$\frac{\text{MrgD+Mrg}^*}{\text{Mrg}^*}$
<i>rMrgA</i>	N.D.	N.D.	N.D.	N.D.	91.5 ± 1.1	100 ± 0
<i>rMrgB5</i>	21.9 ± 18.1	100 ± 0	2.4 ± 2.4	1.9 ± 1.9	N.D.	N.D.
<i>rMrgC</i>	22.0 ± 2.2	26.8 ± 7.9	N.D.	N.D.	1.7 ± 1.7	3.0 ± 3.0
<i>rMrgD</i>	0.9 ± 0.9	0.6 ± 0.6	5.5 ± 5.5	1.2 ± 1.2	N.D.	N.D.

At least three Sprague–Dawley adult DRGs were analyzed by confocal microscopy to determine the extent of colocalization between probe combinations. N.D., not done.