



Supplementary



**Figure S1.** Taste responses of Scnn1<sup>++/++</sup> and Scnn1<sup>aa/bb</sup> mice to control stimuli after dietary intervention. After 4 weeks fed with sodium-adequate, low, or high salt diet, Scnn1<sup>++/++</sup> and Scnn1<sup>aa/bb</sup> mice were subjected to short-term preference tests using an automated gustometer. To do so, animals were either restricted for 22.5 h with access to 2.0 mL water and 1 g food (attractive restriction conditions, (**A**)) or water-deprived for 22.5 h (aversive restriction conditions, (**B**)). Taste solutions and concentrations were presented in random order. Each data point represents a mean ± STE of 5 s presentations from 10 to 11 animals tested. Statistical testing was based on UNIANOVA and post-hoc analysis using Bonferroni's multiple comparison test. Statistical differences were indicated by different letters between individual groups.



**Figure S2.** Latency to initiate the first lick for different taste stimuli after access to amiloridecontaining water. Scnn<sup>1++/++</sup> and Scnn<sup>1aa/bb</sup> mice receiving sodium-adequate diet had either access to 300  $\mu$ M amiloride-containing water 13 h prior to restriction starting or received water without amiloride. The restriction phase lasted for 22.5 h with access to 2.0 mL water ± 300  $\mu$ M amiloride and 1 g food. Mean latency to the first lick for each stimulus concentration was determined by an automated gustometer presenting different concentrated solutions of sucrose (**A**), monopotassium glutamate with inosine 5′monophosphate (MPG+IMP; **B**), sodium chloride (NaCl; **C**), NaCl with amiloride (NaCl+amiloride; **D**), or bitter and sour stimuli (**E**). Each bar represents the mean ± STE from 10 to 16 animals tested. Statistical testing was based on UNIANOVA and post-hoc analysis using Bonferroni's multiple comparison test. Different letters indicate statistical significance.

**Table S1.** Relative expression of ENaC subunits in Scnn1<sup>++/++</sup> and Scnn1<sup>aa/bb</sup> mice. Data represent the relative expression of ENaC subunits normalized to  $\beta$ -actin and eEf2 in isolated taste buds and non-gustatory tissue of Scnn1<sup>++/++</sup> (n = 4) and Scnn1<sup>aa/bb</sup> (n = 6) mice fed with a sodium-adequate diet. Mean variances between Scnn1<sup>++/++</sup> and Scnn1<sup>aa/bb</sup> animals are indicated in %. Statistical testing was based on Student's *t*-test. Differences were considered to be significant if p < 0.05, as indicated in bold.

	ENaC subunit	Scnn1++/++ [33] (mean ± STABW)	Scnn1ªª/bb (mean ± STABW)	Scnn1 <sup>aa/bb</sup> / Scnn1 <sup>++/++</sup> (%)	<i>p-</i> Value
	α	$0.0102 \pm 0.0015$	$0.0151 \pm 0.0053$	148	0.071
fuP	β	$0.0025 \pm 0.0009$	$0.0062 \pm 0.0030$	247	0.049
	γ	$0.0044 \pm 0.0018$	$0.0093 \pm 0.0026$	212	0.011
	α	$0.0390 \pm 0.0054$	$0.0606 \pm 0.0111$	155	0.007
CV+foP	β	$0.0015 \pm 0.0002$	$0.0027 \pm 0.0007$	174	0.011
	γ	$0.0012 \pm 0.0002$	$0.0020 \pm 0.0005$	166	0.015
non-gustatory epithelium	α	$0.0208 \pm 0.0032$	$0.0228 \pm 0.0061$	110	0.567
	β	$0.0007 \pm 0.0002$	$0.0011 \pm 0.0004$	147	0.150
	γ	$0.0008 \pm 0.0001$	$0.0008 \pm 0.0001$	95.0	0.623
kidney	α	$0.0341 \pm 0.0005$	$0.0698 \pm 0.0235$	204	0.018
	β	$0.0157 \pm 0.0026$	$0.0260 \pm 0.0086$	166	0.052
	γ	$0.0278 \pm 0.0145$	$0.0425 \pm 0.0112$	153	0.107
distal colon	α	$0.0268 \pm 0.0085$	$0.0357 \pm 0.0104$	133	0.195
	β	$0.0026 \pm 0.0013$	$0.0051 \pm 0.0017$	193	0.045
	γ	$0.0052 \pm 0.0054$	$0.0091 \pm 0.0068$	173	0.374

**Table S2.** Statistical significance of different factors on the short-term preference tests of Scnn1<sup>++/++</sup> and Scnn1<sup>aa/bb</sup> animals after dietary intervention. After 4 weeks fed with sodium-adequate, low, or high salt diet, 10 to 11 Scnn1<sup>++/++</sup> and Scnn1<sup>aa/bb</sup> mice were subjected to short-term preference tests using an automated gustometer. To do so, animals were either restricted for 22.5 h with access to 2.0 mL water and 1 g food (attractive restriction conditions) or water-deprived for 22.5 h (aversive restriction conditions). Statistical significance (*p*-value) of lick responses to different concentrations of taste solutions based on diet (sodium-adequate, low, or high), genotype (Scnn1<sup>++/++</sup> versus Scnn1<sup>aa/bb</sup>), and diet X genotype interactions are shown. Statistical testing was based on UNIANOVA and post-hoc analysis using Bonferroni's multiple comparison test. Differences were considered to be statistically significant if *p* < 0.05, as indicated in bold.

Protocol	Substance	Concentration	Diet	Genotype	Diet X genotype
		10	0.736	0.761	0.609
		30	0.022	0.021	<0.001
	sucrose	100	0.182	0.412	0.376
		300	0.461	0.784	0.842
		1000	0.479	0.176	0.151
		1	0.282	<0.001	<0.001
		3	0.013	0.032	0.002
	MPG+IMP	10	0.957	<0.001	<0.001
		30	0.914	0.824	0.962
		100	0.166	0.985	0.545
		10	0.140	0.214	0.089
attractivo		30	0.009	0.122	0.014
attractive	NaCl	100	0.739	<0.001	0.029
		300	0.303	0.034	0.042
		1000	0.359	0.004	0.049
		10	0.250	0.058	0.125
	NaCl+amilorid e	30	0.599	0.127	0.431
		100	0.352	0.204	0.297
		300	0.430	0.078	0.205
		1000	0.596	0.490	0.548
	amiloride	0.1	0.414	0.001	0.012
	IMP	0.1	0.380	0.457	0.724
	denatonium	1	0.293	0.663	0.764
	citric aid	100	0.113	0.006	0.009
aversive		0.1	0.460	0.002	0.006
		0.3	0.545	0.001	0.035
	denatonium	1	0.308	0.030	0.038
		3	0.893	0.256	0.662
		10	0.914	0.240	0.884
		1	0.124	<0.001	0.003

Protocol	Substance	Concentration	Diet	Genotype	Diet X genotype
	citric acid	3	0.380	0.963	0.364
		10	0.362	<0.001	<0.001
		30	0.209	0.045	0.105
		100	0.005	0.022	0.003
		10	0.171	0.081	0.090
		30	0.861	0.632	0.759
	NaCl	100	0.183	0.505	0.525
		300	0.002	0.050	<0.001
		1000	<0.001	<0.001	<0.001
	NaCl+amilorid e	10	0.068	0.754	0.035
		30	0.002	0.144	<0.001
		100	0.067	0.307	0.284
		300	0.002	0.354	0.007
		1000	0.001	0.001	<0.001
	amiloride	0.1	0.389	0.002	0.028
	sucrose	300	0.009	0.815	0.007
	sucralose	10	0.080	0.166	0.025
	MPG+IMP	100	0.001	0.004	<0.001

**Table S3.** Statistical significance of different factors on the short-term preference tests of Scnn1<sup>++/++</sup> and Scnn1<sup>aa/bb</sup> animals. After 4 weeks fed with sodium-adequate, low, or high salt diet, 10 to 11 Scnn1<sup>++/++</sup> and Scnn1<sup>aa/bb</sup> mice were subjected to short-term preference tests using an automated gustometer. To do so, animals were either restricted for 22.5 h with access to 2.0 mL water and 1 g food (attractive restriction conditions) or water-deprived for 22.5 h (aversive restriction conditions). Data represent the statistical significance (*p*-value) of diet, genotype, concentration, and a different combination of them, based on all tested concentrations of one substance (first 4 listed substances for each protocol were tested for 5 concentrations, whereas the remaining were only checked for 1 concentration). *p*-Values were based on UNIANOVA and post-hoc analysis using Bonferroni's multiple comparison test. Differences were considered to be statistically significant if *p* < 0.05, as indicated in bold.

Protocol	Substance	Substance Factor	
		diet	0.797
		concentration	<0.001
	sucrose	diet X concentration	0.134
		genotype	0.743
		genotype X concentration	0.181
		diet X genotype	0.889
		diet X genotype X concentration	0.034
		diet	0.918
		concentration	<0.001
		diet X concentration	0.119
	MPG+IMP	genotype	0.001
		genotype X concentration	0.005
attractive		diet X genotype	0.031
attractive		diet X genotype X concentration	0.026
	NaCl	diet	0.360
		concentration	0.030
		diet X concentration	0.005
		genotype	<0.001
		genotype X concentration	0.640
		diet X genotype	0.001
		diet X genotype X concentration	0.004
		diet	0.075
		concentration	0.067
	NaCl+amiloride	diet X concentration	0.871
		genotype	0.001
		genotype X concentration	0.522

Protocol	Substance	Factor	<i>p</i> -Value
		diet X genotype	<0.001
		diet X genotype X concentration	0.974
		diet	0.414
	amiloride	genotype	0.001
		diet X genotype	0.012
		diet	0.380
	IMP	genotype	0.457
		diet X genotype	0.724
		diet	0.293
	denatonium	genotype	0.663
		diet X genotype	0.764
		diet	0.113
	citric aid	genotype	0.006
		diet X genotype	0.009
		diet	0.089
		concentration	<0.001
	denatonium	diet X concentration	0.757
		genotype	<0.001
		genotype X concentration	<0.001
		diet X genotype	<0.001
		diet X genotype X concentration	0.002
		diet	0.015
		concentration	<0.001
	citric acid	diet X concentration	0.497
		genotype	<0.001
aversive		genotype X concentration	<0.001
		diet X genotype	<0.001
		diet X genotype X concentration	<0.001
		diet	<0.001
		concentration	<0.001
		diet X concentration	0.012
	NaCl	genotype	0.060
		genotype X concentration	0.004
		diet X genotype	<0.001
		diet X genotype X concentration	<0.001
	NaCl+amiloride	diet	<0.001
		concentration	<0.001

Protocol	Substance	Factor	<i>p</i> -Value
		diet X concentration	<0.001
		genotype	0.758
		genotype X concentration	0.132
		diet X genotype	0.002
		diet X genotype X concentration	<0.001
		diet	0.389
	amiloride	genotype	0.002
		diet X genotype	0.028
		diet	0.009
	sucrose	genotype	0.815
		diet X genotype	0.007
		diet	0.080
	sucralose	genotype	0.166
		diet X genotype	0.025
		diet	0.001
	MPG+IMP	genotype	0.004
		diet X genotype	<0.001

**Table S4.** Statistical significance of different factors on the short-term preference tests of Scnn1<sup>++/++</sup> and Scnn1<sup>aa/bb</sup> animals after amiloride intervention for 36 h. After receiving sodium-adequate diet, animals either had access to water or amiloride-containing water (300  $\mu$ M), following short-term preference testing in an automated gustometer. Data represent the statistical significance (*p*-value) of an intervention (water with or without amiloride), genotype (Scnn1<sup>++/++</sup> versus Scnn1<sup>aa/bb</sup>), and intervention X genotype on different taste solutions and their concentrations. Statistical testing was based on UNIANOVA and post-hoc analysis using Bonferroni's multiple comparison test for data points of 10 to 16 animals. Differences were considered to be statistically significant if *p* < 0.05, as indicated in bold.

Substance	Concentration	Intervention	Genotype	Intervention X genotype
	10	0.002	0.160	0.001
sucrose	100	0.007	0.427	0.002
	1000	<0.001	0.776	<0.001
	1	0.633	0.531	0.198
MPG+IMP	10	0.001	0.003	<0.001
	100	<0.001	0.036	<0.001
	10	0.008	0.162	0.033
NaCl	100	<0.001	0.001	<0.001
	1000	<0.001	0.011	<0.001
NaCl+amiloride	10	<0.001	0.019	<0.001
	100	<0.001	0.525	<0.001
	1000	0.002	0.030	0.002
denatonium	10	0.583	0.599	0.684
citric aid	100	0.006	0.488	0.026

**Table S5.** Statistical significance of different factors on the short-term preference tests of Scnn1<sup>++/++</sup> and Scnn1<sup>aa/bb</sup> animals after access to amiloride-containing drinking water for 36 h. After receiving sodium-adequate diet, animals either had access to water or amiloride-containing water (300  $\mu$ M), following short-term preference testing in an automated gustometer. Data represent the statistical significance (*p*-value) of intervention (water with or without amiloride), genotype, concentration, and different combination/interaction of them based on all 5 concentrations tested for 1 stimulus/substance (for denatonium and citric acid only 1 concentration was tested). Test was based on UNIANOVA and post-hoc analysis using Bonferroni's multiple comparison test relying on data of 10 to 16 animals. Differences were considered to be statistically significant if *p* < 0.05, as indicated in bold.

Substance	Factor	<i>v</i> -Value
	intervention	<0.001
	concentration	<0.001
	intervention X concentration	<0.001
sucrose	genotype	0.998
	genotype X concentration	0.564
	intervention X genotype	<0.001
	intervention X genotype X concentration	<0.001
	intervention	<0.001
	concentration	<0.001
	intervention X concentration	<0.001
MPG+IMP	genotype	0.001
	genotype X concentration	0.119
	intervention X genotype	<0.001
	intervention X genotype X concentration	0.001
	intervention	<0.001
	concentration	<0.001
	intervention X concentration	0.003
NaCl	genotype	<0.001
	genotype X concentration	0.019
	intervention X genotype	<0.001
	intervention X genotype X concentration	0.001
	intervention	<0.001
	concentration	<0.001
	intervention X concentration	0.002
NaCl+amiloride	genotype	0.007
	genotype X concentration	0.364
	intervention X genotype	<0.001
	intervention X genotype X concentration	0.014
	intervention	0.583
denatonium	genotype	0.599
	intervention X genotype	0.684
	intervention	0.006
citric aid	genotype	0.488
	intervention X genotype	0.026