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Electrophysiological characterization of a new member of the RCK family of rat brain K^+ channels

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A novel member of the RCK family of rat brain K+ channels, called RCKZ, has been sequenced and expressed in *Xenopus* oocytes. The K+ currents were voltage-dependent, activated within 20 ms (at 0 mV), did not inactivate in 5 s, and had a single channel conductance in frog Ringers of 8.2 pS. Compared to other members of the RCK family the pharmacological profile of RCK2 was unique in that the channel was resistant to block $(IC_{50} = 3.3 \,\mu M)$ by charybdotoxin [(1988) Proc. Natl. Acad. Sci. USA 85, 3329-3333] but relatively sensitive to 4-aminopyridine (0.3 mM), tetraethylammonium (1.7 mM), a-dendrotoxin (25 nM), noxiustoxin (200 nM), and mast cell degranulating peptide (200 nM). Thus, RCK2 is a non-inactivating delayed rectifier K⁺ channel with interesting pharmacological properties.

Potassium channel; cDNA cloning; cDNA expression; Delayed rectifier; *Xenopus* oocyte

proteins which regulate the transmembrane diffusion of transmitter release in rat brain synaptosomes [1].
K⁺ ions. Channel activation controls neuronal ex-
RCK2, a cDNA encoding a rat brain K⁺ channel has K^+ ions. Channel activation controls neuronal ex-
citability through repolarization of the action potential and modulation of the frequency of repetitive firing [5]. clone reported here encodes a polypeptide which differs
From whole-cell and single channel electrophysiological from RCK2 at only one residue and hence will be refe From whole-cell and single channel electrophysiological from RCK2 at only one residue and hence will be refer-
measurements [25.27], neurons are thought to express red to as RCK2, also. Our electrophysiological results measurements [25,27], neurons are thought to express red to as RCK2, also. Our electrophysiological results in
heterogeneous populations of K⁺ channels which differ are closely comparable to those obtained by Grupe et al heterogeneous populations of K⁺ channels which differ are closely comparable to those obtained by Grupe et al.
in their biophysical and pharmacological properties. [10], except that in our experiments charybdotoxin in their biophysical and pharmacological properties. Until recently the structural basis of K⁺ channel diversi-
ty was not known; however, molecular cloning methods rent than was reported by Grupe et al. [10]. ty was not known; however, molecular cloning methods have now shown that mammalian brain mRNA encodes several distinct voltage-activated K^+ channels [2,8,15, several distinct voltage-activated K channels $[2,0,15]$, 2. MATERIALS AND METHODS 22,24,25]. Expression of cloned rat brain K⁺ channels provides a new basis for understanding the relationships between primary structure and channel function. 2.1. *Isolation and sequencing of cDNA*

Using an oligonucleotide probe encoding a strictly conserved sequence located upstream of the membrane the amino acid sequence Asn-Glu-Tyr-Phe-Asp-Arg (position $\frac{1}{2}$ channels we $\frac{1}{2}$ 82-88 of RCK2) which is conserved in most of the known voltagespanning core region of several known K^+ channels, we have isolated a cDNA clone which encodes a new member of the RCK family of K^+ channels $[2,10]$. When expressed in *Xenopus* oocytes this clone produces non-inactivating delayed rectifier K^+ current. Its sen-

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1, INTRODUCTION sitivity to blockers such as 4-aminopyridine, dendrotoxin and mast cell degranulating protein suggests that it Voltage-activated K⁺ channels are integral membrane may be related to the K⁺ channels which affect neuro-
oteins which regulate the transmembrane diffusion of transmitter release in rat brain synaptosomes [1].

recently been described by Grupe et al. *[10]*. The cDNA clone reported here encodes a polypeptide which differs

Rat brain cDNA libraries enriched for full-length inserts [7] were screened at low stringency using an oligonucleotide probe encoding
the amino acid sequence Asn-Glu-Tyr-Phe-Phe-Asp-Arg (position activated K+ channels. Run-off RNA transcripts of positive clones were made as described previously [12] and stage V-VI Xenopus oocytes were injected with 10 ng of RNA in 75 nl 0.1 M KC1 and screened for K^+ current expression. RNA from one clone (RCK2) bearing an approximately 5 kb insert, gave large K^+ currents, and had a DNA sequence which was different from all known K' channels.

2.2. *Electrophysiological recording Correspondence address: G.E. Kirsch, Baylor College of Medicine,* Cocytes were incubated at 19°C in modified Barth's solution for
Dept. of Anesthesiology, One Baylor Plaza, Houston, TX 77030, 2-7 days and then tested for voltage clamp [12]. Manually defolliculated oocytes were placed in a recording chamber continuously superfused at 3 ml/min with a test solution consisting of (mM): 120 NaOH, 120 methanesulfonic acid, 2.5 KCI, and 10 Hepes, adjusted to pH 7.3 with NaOH. Occytes were impaled with 3 M KCI-filled micropipettes (resistance 1-2 MO).

 $\hat{\psi}$ $\begin{array}{c} \frac{1}{2} \\ \frac{1}{2} \end{array}$

Fig. 1. Amino acid sequence alignment of members of the RCK family. Identical amino acids are indicated by dashed line. Gaps (dotted lines) have been introduced to achieve maximum homology. Putative transmembrane segments (Si-S6) are indicated by solid bars. RCK-1, -3, -4 and -5 are from Baumann et al. [2] and Stühmer et al. [22]. Sequence alignment was obtained using EuGene software (Molecular Biology Information Resources, Department of Cell Biology, Baylor College of Medicine).

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4-Aminopyridine (4-AP), tetraethylammonium chloride (TEA) and 'mast cell degrahulating peptide (MCDP) were obtained from Sigma Chemical Co. (S:. Louis, MO). Charybdotoxin (ChTX) was obtained from Latoxan (Rosans, France) and is prepared by the method of Gimenez-Gallego [9]. Noxiustoxin (NTX) and α -dendrotoxin (DTX) were the generous gifts of Dr L.D. Possani and Dr M.P. Blaustein, respectively.

Single channel recording was performed in oocytes dissected free of the vitelline envelope and patch clamped (121 using fire-polished, Sylgard-coated micropipettes of 2-5 M Ω resistance when filled with the test solution described above. Data acquisition and analysis was performed using pCLAMP software (Axon Instruments, Burlingame, CA). Single channel records were filtered at 500-1000 Hz and digitized at 1000-2000 Hz. Where appropriate, data are expressed as $mean \pm SE$.

3. RESULTS

3,1. *Structural properties of ROY.2*

Putative $K⁺$ channel clones were isolated by hybridization screening of rat brain cDNA libraries with a K^+ channel-specific oligonucleotide probe. Expression screening yielded a clone whose sequence (Fig. 1) and electrophysiological properties clearly place it in the

Fig. 2. Steady-state current/voltage relationship of RCK2. Oocytes injected with 10 ng mRNA transcript were voltage-clamped using two intracellular microelectrodes (A) and stimulated with a series of test pulses of -50 to $+50$ mV in 10 mV increments from a holding potential of -80 mV. Linear leakage and capacitative currents were substracted digitally using a P/4 substraction protocol. Panel (A) shows an $I-V$ family obtained with 4.7 s pulses. Panel (B) shows the ensemble average of 100 test pulses to 0 mV in a cell-attached membrane patch which contained a single channel. The vertical calibration is units of probability of opening. In panel (C) steady-state currents from microelectrode measurements were converted to chord conductances using a reversal potential of -92 mV (measured in separate experiments), and fitted to Boltzmann distributions 'with midpoint of slope factor (mV): -11.5 and 12.2 for RCK2. Conductances were normalized to the maximum estimated from the Boltzmann fit.

RCK family originally described by Baumann et al. [2] and its deduced amino acid sequence (Fig. 1) shows that it is nearly identical to RCK2, recently described by Grupe et al. [IO] and KV2, recently described by Swanson et al. $[23]$. A single amino acid (leucine-241) is replaced by a serine in the previous reports [10,23].

Hydropathy analysis indicates six hydrophobic, putative membrane-spanning regions (S1–S6). This core region of the molecule shows strong sequence homology with all members of the RCK family [2,22]. Highest levels of sequence identity (up to 89%) were observed from the beginning of the S4 to the end of the S6 regions. Less conservation is evident in the terminal regions and the putative extracellular 51-52 and S3-S4 linker regions which are longer in RCK2 than in the other RCK variants. These differences may account for some of the unique functional characteristics of RCK2 described below.

3.2. *Electrophysiological properties of RCK.2*

whereas under the same experimental conditions, outuninjected oocytes. The expression of exogenous mRNA therefore is responsible for $>90\%$ of the outward current recorded in the injected oocytes. Panel A shows an I-V family of superimposed currents evoked by test pulses ranging from -50 to $+50$ mV, in 10 mV increments from a holding potential of -80 mV. The records are corrected for linear capacitative and leakage currents, and therefore represent the activation of a voltage-dependent conductance. Fig. 2A shows that The steady-state current/voltage (I-V) relationship for RCK2 is shown in Fig. 2. Voltage-dependent outward currents of 2–6 μ A (at a test potential of +50 mV) were recorded in oocytes injected with 10 ng RNA, ward. currents of less than 150 nA were recorded in RCKZ expresses an outward current which does not inactivate during 4.7 s pulses. A similar lack of inactivation is characteristic of RCKl and RCK5; in contrast, RCK3 and RCK4 inactivate 80-100% over this time period [22]. We have not looked for ultra-slow components of inactivation which might become apparent when the holding potential is made more positive. It is clear, however, that RCKZ lacks fast inactivation.

In order to resolve the time course of activation, we rccnrded single channel currents in cell-attached membrane patches. Fig. 2B shows a typical recording of the ensemble average response evoked by 100 test pulses to 0 mV in a membrane patch containing only one channel. Activation was complete within about 15 ms. The average rise time (90% of peak) was 12.4 ± 2.2 ms ($n = 8$ patches) at 0 mV, a value within the range of activation times measured in other members of the RGK family [10,22].

Fig. 2C shows the steady-state voltage dependence of activation. The conductance/voltage $(G-V)$ relationship was fitted by a Boltzmann distribution with

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average midpoint and slope factor of -12.3 ± 4.3 and average midpoint and slope factor of -12.3 ± 4.3 and $A \uparrow \uparrow$
11.9±1.7 mV (n = 6), respectively. In the other $A \uparrow \uparrow$ members of the RCK family midpoints range from -22 to -34 mV and slope factors range from 5 to 17 mV [22]; RCK2 therefore has a relatively high threshold for activation compared with other members of the RCK family [IO,22].

Fig. 3 demonstrates the $[K^+]$ -dependence of RCK2 currents. Reversal potentials were measured from tail currents evoked by a double pulse protocol shown in Fig. 3A. Conditioning pulses to 0 mV activated out-

Fig. 3. Effects of changing extracellular $[K^+]$ on RCK2. Na⁺ was replaced by the desired amount of K^+ . Tail currents were measured by the pulse protocol shown in panel (A). Lower set of superimposed traces in panel (A) show tail currents measured in 100 mM K+. In panel (B) peak tail currents are plotted as a function of test pulse potential, in 40 (circles) and 100 (triangles) mM K+. Zero current potentials were measured by interpolation of the isochronal $I-V$ curves and pooled daia (linear axis) from 4 oocytes and plotted as a function of $[K^+]_0$ (logarithmic axis) in panel (C). Data are plotted as $mean ± SD$. The straight line was fitted by least squares regression.

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Fig. 4. Single channel RCKZ currents. Cell-attached patch clamp of RCK2-injected oocytes revealed a voltage-activated outward current which was absent in uninjected oocytes. Test pulses of varying amplitude were delivered repetitively at 0.5 Hz from a holding potential of -80 mV. Typical records of single channel activity are shown in (A) (low-pass filtered at 1000 Hz). Panel (B) shows the first 200 ms of the first trace in panel (A) at a faster time base. Records were idealized, amplitude hisstograms (not shown) and open time histograms(C) were constructed. Mean single channel amplitudes obtained by fitting the histograms to single Gaussian distributions and mean open times were obtained by fitting single exponential decay functions. Panel (D) shows unitary $I-V$ relationship for pooled data from 8 patches. The straight line is a least squares fit of the data.

c c mV were used to assess the fully activated conductance. **0** Isochronal tail currents in 100 (triangles) and 40 (circles) mM [K⁺]_o are plotted as a function of test pulse potential in Fig. 3B. Reducing $[K^+]_0$ from 100 to 40 mM shifted the zero current potential by -30 mV. The pooled data from 5 experiments (Fig. 3C) show that the conductance expressed by RCK2 is highly selective for K^+ over Na⁺ or Cl⁻. The 54.5 mV/decade slope of the decade range measured in other RCK variants [22].

> Single channel currents were measured in eight cellattached patches. Representative records are shown in Fig. 4A,B. As shown in Fig. 4A channel activation evoked by test pulses to 0 mV occurred after a. brief latency in long bursts which often lasted the entire duration of the test pulse. This feature can account for lack of inactivation in whole-cell RCK2 currents. At a faster time base (Fig. 4B) bursts were found to consist of long openings separated by brief $(< 1$ ms) closed intervals. Closure of the channel to a subconductance state is also

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illustrated in this record. The main conductance state quite closely with the work of Grupe et al. [10] who first had a mean open time of 12 ms (Fig. 4C) and mean described expression of RCK2 in oocytes. Both reports amplitude of 0.89 pA. Single channel conductance show that RCK2 codes for a K⁺ channel which activates estimated from pooled data in Fig. 4C was 8.2 pS $(-10 - 10)$ slowly (tens of milliseconds), has no inactivation in th estimated from pooled data in Fig. 4C was 8.2 pS (-10 slowly (tens of milliseconds), has no inactivation in the to $+60$ mV), a value which is within the 4.7-10.2 pS range 500-5000 ms, and has sensitivity to several to $+60$ mV), a value which is within the 4.7–10.2 pS range 500–5000 ms, and has sensitivity to several range of conductances of other RCK variants $[10.22]$; K^+ -channel blockers including DTX. MCDP, TEA and range of conductances of other RCK variants [10,22]; K^+ -channel blockers including DTX, MCDP, TEA and and mean open time was 13.9 ± 1.3 ms ($n = 8$ patches). $4-AP$. However, in our work and that of Swanson et al.

summarized in Table I. Block was measured at one or sequences of RCK2 reported by Grupe et al. [10] and more concentrations and the ratio of test/control cur-
that of the ChTX-insensitive KV2 channel [23] are idenrents evoked by test pulses to 10 mV, from a holding tical. A plausible explanation is that different toxins potential of -80 mV was used to estimate the IC_{50} , were used. Our sample of ChTX was prepared by the potential of -80 mV was used to estimate the IC_{50} , were used. Our sample of ChTX was prepared by the assuming single site binding [20]. RCK2 shows high sen-
method of Gimenez-Gallego et al. [9] whereas that used assuming single site binding [20]. RCK2 shows high sen-
sitivity to all the blocking agents tested except ChTX. by Grupe et al. [10] was prepared by the method of Our sample of ChTX blocked outward current $(IC_{50} \approx$ Miller et al. [16]. Swanson et al. [23] obtained toxins 30 nM) in oocytes injected with another K^+ channel prepared by both methods and it is unclear which toxin clone which is closely homologous to the ChTX- was tested on their RCK2 channel. When both toxin sensitive RCK1 channel. Thus, the ChTX insensitivity preparations were tested on delayed rectifier currents in
of RCK2 is a real feature of the channel rather than an lymphocytes [3] the blocking potency of the latter artifact of inactive toxin. Interestingly, RCK2 is sen-
sitive to noxiustoxin (NTX), a scorpion venom-derived Only the polypeptide isolated by Gimenez-Gallego et al. sitive to noxiustoxin (NTX), a scorpion venom-derived Only the polypeptide isolated by Gimenez-Gallego et al.
peptide toxin which is related to ChTX by sequence [9] has been sequenced and it is known that Leirus quinpeptide toxin which is related to ChTX by sequence [9] has been sequenced and it is known that *Leirus quin*-
auestriatus venom contains at least two closely related

with roughly the same potency as block of RCK1, as block of RCK5 [22]. RCK2 is sensitive to block by less discriminating. The use of synthetic toxins of TEA (IC₅₀ = 1.7 mM), consistent with the notion that known amino acid sequence may be necessary to resolve delayed rectifiers are selectively blocked by this drug. this issue. However, as shown in Table I, RCK2 is also quite sen-
sitive to 4-AP, which is often considered a selective of the critical amino acids for binding of K^+ channel sitive to 4-AP, which is often considered a selective of the critical amino acids for binding of K^+ channel
blocker of I_A , a transient K^+ current. RCK2 appears to peptide toxins other than ChTX (prepared by the blocker of I_A , a transient K⁺ current. RCK2 appears to be slightly more sensitive to 4-AP than the other RCK

described identical rat brain cDNA clones named RCK2 clones [18], and antibodies to synthetic polypeptides [10] and $K\sqrt{2}$ [23], both of which encode a rat brain K^+ deduced from the mouse homolog of RCK1 recognize [10] and K_v2 [23], both of which encode a rat brain K⁺ deduced from the mouse homolog of RCK1 recognize channel. Our results and those of Swanson et al. [23] the purified rat brain DTX receptor [19]. In neurons agree quite closely, particularly with regard to ChTX these three toxins all interact allosterically with one
insensitivity. Our electrophysiological results agree another, suggesting that although they do not occupy

4-AP. However, in our work and that of Swanson et al. [23], RCK2 was relatively insensitive to block by ChTX. 3.3. *RCK2 pharmacology* **This difference cannot arise from structural differences**
We tested RCK2 sensitivity to K⁺ channel blockers as in the expressed channels since the deduced amino acid We tested RCK2 sensitivity to K^+ channel blockers as in the expressed channels since the deduced amino acid
summarized in Table I. Block was measured at one or sequences of RCK2 reported by Grupe et al. [10] and that of the ChTX-insensitive KV2 channel [23] are idenby Grupe et al. [10] was prepared by the method of lymphocytes [3] the blocking potency of the latter mologies [26].
RCK2 is sensitive to block by a dentrotoxin (DTX) OhTX isoforms [13]. We suggest, therefore, that RCK2 ChTX isoforms [13]. We suggest, therefore, that RCK2 may be sensitive to only one of the isoforms, whereas whereas MCDP blocked RCK2 with the same potency other members of the RCK family (e.g. RCK1) may be known amino acid sequence may be necessary to resolve

be slightly more sensitive to 4-AP than the other RCK method of Miller et al. [16] which is thought to bind to variants (range 1-13 mM) [22]. a site located on the extracellular linker between transmembrane segments $S5-S6$ [14]. It is known, 4. DISCUSSION however, that MCDP, DTX and ChTX all bind to protein receptors that, when deglycosylated, have a After this paper was submitted two other groups molecular weight similar to that estimated from RCK
described identical rat brain cDNA clones named RCK2 clones [18], and antibodies to synthetic polypeptides the purified rat brain DTX receptor [19]. In neurons another, suggesting that although they do not occupy the same site on the channel, their binding sites may be close together. One possibility is that all of the toxin sites are on the S5-S6 linker. In that case the MCDP-Effect of K⁺ channel blockers on RCK2-induced current and DTX-resistance of both RCK3 and RCK4 may be due to subtle alterations in the variable regions of the $S5-86$ linker. A possible site is located at RCK1 position 353 which is occupied by a negatively charged amino acid in RCKl, RCKS and RCK2 but is occupied by either Ser or Thr in RCK3 and RCK4. Interestingly, in the latter two clones, position 352 is occupied by a helixdistorting Pro.

^a Rel. block = $1-(R_{\star,dm}/R_{\star,\text{control}})$ Studies of the functional characteristics of muta-

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 K^+ channel variants [22], have helped to identify the $33-38$.
molecular machinery responsible for voltage-dependent [8] Frech, G.C., VanDongen, A.M.J., Schuster, G., Brown, A.M. molecular machinery responsible for voltage-dependent [8] Frech, G.C., VanDongen, A.M.J., Schuster, and Joho, R.H. (1989) Nature 340, 642–645. gating. In both K^+ channels and Na^t channels rapid, voltage-dependent inactivation is thought to be localiz ed in intracellular loops which connect the adjacent subunits or pseudosubunit repeats in $Na⁺$ channels $[4,21]$. As noted by Stühmer et al. $[22]$, among the members of the RCK family, fast inactivation is present only jn the RCK4 variant which also has a long aminoterminus. RCK2, with its relatively short N-terminus and lack of inactivation, conforms to this principle. Site-directed mutagenesis should enable us to test the structure/function correlations suggested by comparisons between different delayed rectifier K⁺ chan-FILM 1988 1. SL.V 1151 McKinnon, D. (1989) J. Biol. Chem. 264, 8230-8236.

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