

- liams, S., McDermid, H., Dumanski, J. P., Biegel, J., Bell, C. J., and Emanuel, B. S. (1996). Regional localization of over 300 loci on human chromosome 22 using a somatic cell hybrid mapping panel. *Genomics* **35**: 275–288.
3. Cox, D. R., Burmeister, M., Price, E. R., Kim, S., and Myers, R. M. (1990). Radiation hybrid mapping: A somatic cell genetic method for constructing high resolution maps of mammalian chromosomes. *Science* **250**: 245–250.
 4. Dutrillaux, B., and Viegas-Pequignot, E. (1985). High resolution R- and G-banding in the same preparation. *Hum. Genet.* **57**: 93–95.
 5. Engelen, J. J. M., de Die-Smulders, C. E. M., Fryns, J. P., Hoovers, J. M. N., Albrechts, J. C. M., Loots, W. J. G., Jacobs, M. E., and Hamers, A. J. H. (1994). Partial trisomy and monosomy 8p due to inversion duplication. *Clin. Genet.* **45**: 203–207.
 6. Frid, M. G., Shekhonin, B. V., Koteliansky, V. E., and Glukhova, M. A. (1992). Phenotypic changes of human smooth muscle cells during development: Late expression of heavy caldesmon and calponin. *Dev. Biol.* **153**: 185–193.
 7. Glukhova, M. A., Kabakov, A. E., Belkin, A. M., Frid, M. G., Ornaty, O. I., Zhidkova, N. I., and Koteliansky, V. E. (1986). Meta-vinculin distribution in adult human tissues and cultured cells. *FEBS Lett.* **207**: 139–141.
 8. Kurahashi, H., Akagi, K., Karakawa, K., Nakamura, T., Dumanski, J. P., Sano, T., Okada, S., Takai, S., and Nishisho, I. (1994). Isolation and mapping of cosmid markers on chromosome 22 including one with the submicroscopically deleted region of DiGeorge syndrome. *Hum. Genet.* **93**: 248–254.
 9. Mulder, M. P., Wilke, M., Langeveld, A., Wilming, L. G., Hagemeyer, A., Van Drunen, E., Zwarthoff, E. C., Riegman, P. H. J., Deelen, W. H., Van den Ouweland, A. M. W., Halley, D. J. J., and Meijers, C. (1995). Positional mapping of loci in the DiGeorge critical region at chromosome 22q11 using a new marker (D22S1 83). *Hum. Genet.* **96**: 133–141.
 10. Owens, G. K. (1995). Regulation of differentiation of vascular smooth muscle cells. *Physiol. Rev.* **75**: 487–517.
 11. Sambrook, J., Fritsch, E. F., and Maniatis, T. (1989). "Molecular Cloning: A Laboratory Manual," Cold Spring Harbor Laboratory Press, Cold Spring Harbor, New York.
 12. Schuler, G. D., Boguski, M. S., Steward, E. A., Stein, L. D., Gypay, G., Rice, K., White, R. E., Rodriguez-Tom, P., Aggarwal, A., Bajorek, E., Bentolila, S., Birren, B. B., Butler, A., Castle, A. B., Chiannilkulchai, N., Chu, A., Clee, C., Cowles, S., Day, P. J. R., Dibling, T., Drouot, N., Dunham, I., Duprat, S., East, C., Edwards, C., Fan, J.-B., Fang, N., Fizames, C., Garret, C., Green, L., Haley, D., Harris, M., Harrison, P., Brady, S., Hicks, A., Holloway, E., Hui, L., Hussain, S., Louis-Dit-Sully, C., Ma, J., MacGilvery, A., Mader, C., Maratukulam, A., Matise, T. C., McKusick, K. B., Morissette, J., Mungall, A., Muselet, D., Nusbaum, H. C., Page, D. C., Peck, A., Perkins, S., Piercy, M., Qin, F., Quackenbush, J., Ranby, S., Reif, T., Rozen, S., Sanders, C., She, X., Silva, J., Slonim, D. K., Soderlund, C., Sun, W.-L., Tabar, P., Thangarajah, T., Vega-Czarny, N., Vollrath, D., Voyticky, S., Wilmer, T., Wu, X., Adams, M. D., Auffray, C., Walter, N. A. R., Brandon, R., Dehejia, A., Goodfellow, R., Houlgatte, R., Hudson, J. R., Jr., Ide, S. E., Iorio, K. R., Lee, W. Y., Seki, N., Nagase, T., Ishikawa, K., Nomura, N., Phillips, C., Polymeropoulos, M. H., Sanduski, M., Schmitt, K., Berry, R., Swanson, K., Torres, R., Venter, J. C., Sikela, J. M., Beckmann, J. S., Weissenbach, J., Myers, R. M., Cox, D. R., James, M. R., Bentley, D., Deloukas, P., Lander, E. S., and Hudson, T. J. (1996). A gene map of the human genome. *Science* **274**: 540–546.
 13. Takahashi, K., Hiwada, K., and Kokubu, T. (1988). Vascular smooth muscle calponin: A novel T-like protein. *Hypertension* **11**: 620–626.
 14. Takeuchi, K., Takahashi, K., Abe, M., Nishida, W., Hiwada, K., Nabeya, T., and Maruyama, K. (1991). Co-localization of immunoreactive forms of calponin with actin cytoskeleton in platelets, fibroblasts and smooth muscle. *J. Biochem.* **109**: 311–316.
 15. Van der Loop, F. T. L., Schaart, G., Timmer, E. D. J., Ramaekers, F. C. S., and Van Eys, G. J. J. M. (1996). Smoothelin, a novel cytoskeletal protein specific for smooth muscle cells. *J. Cell Biol.* **134**: 401–411.
 16. Van der Loop, F. T. L., Gabbiani, G., Kohonen, G., Ramaekers, F.-C. S., and Van Eys, G. J. J. M. (1997). Differentiation of smooth muscle cells in human blood vessels as defined by smoothelin, a novel marker for the contractile phenotype. *Atheroscl. Thromb. Vas. Biol.* **17**: 665–671.
 17. Wiegant, J., and Dauwerse, J. G. (1995). Multiple-coloured chromosomes by in situ hybridization. In "Human Chromosomes: Principles and Techniques" (R. S. Verma and A. Babu, Eds.), 2nd ed., McGraw-Hill, New York.

Localization of the Gene Encoding the Ran-Binding Protein RanBP2 to Human Chromosome 2q11–q13 by Fluorescence *in Situ* Hybridization

Heike Krebber,* Holger Bastians,* Jörg Hoheisel,† Peter Lichter,‡ Herwig Ponstingl,*¹ and Stefan Joos‡

*Division for Molecular Biology of Mitosis, 0230,

‡Organization of Complex Genomes, 0845, and †Molecular-Genetic Genome Analysis, 0846, German Cancer Research Center, Im Neuenheimer Feld 280, D-69120 Heidelberg, Germany

Received March 5, 1997; accepted April 17, 1997

Ran (Ras-related small nuclear protein) is a member of the Ras superfamily of small GTP-binding proteins. It is involved in cell-cycle progression, nucleocytoplasmic transport, and pre-mRNA processing (7). Ran is predominantly located in the nucleus and cycles between the GTP-bound active and the GDP-bound inactive state (1). RanBP2 (Ran-binding protein 2) is the largest protein of the nuclear pore complex, localized to its cytoplasmic fibers (10–12). It contains four Ran-binding domains and a C-terminal cyclophilin-related region (11, 12) that has been found to act as chaperone for red/green opsin (3). A RanBP2-specific antibody inhibits nuclear protein import, indicating a functional role of RanBP2 in this process (12). The RanGTPase-activating protein RanGAP1 is posttranslationally modified by a small ubiquitin-related polypeptide and is thereby targeted to RanBP2 (6).

To assign the chromosomal locus of the RanBP2 gene, fluorescence *in situ* hybridization was performed according to the

¹ To whom correspondence should be addressed at the German Cancer Research Center, Division for Molecular Biology of Mitosis, 0230, Im Neuenheimer Feld 280, D-69120 Heidelberg, Germany. Telephone: (+49-6221-423408). Fax: (+49-6221-423460).

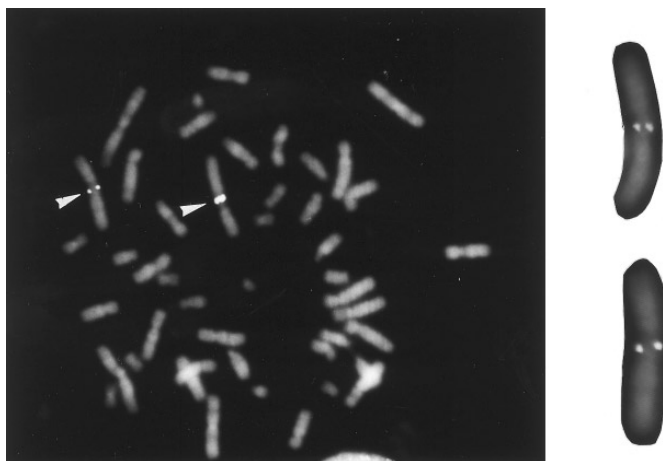


FIG. 1. Localization of the human RanBP2 gene on chromosome 2q11–q13 by fluorescence *in situ* hybridization. Normal male/female chromosomes were hybridized with biotinylated probes for RanBP2. The hybridization site on chromosome 2 is indicated by arrowheads.

protocol described (4). cDNA probes from a domain of RanBP2 (1) were used to identify yeast artificial chromosomes harboring the gene encoding RanBP2. The hybridization procedure was carried out as described (4). A YAC that hybridized with the RanBP2 probe (ICRF-Y-900C04133) was used as a template in amplification of human-specific DNA sequences by interspersed long-range PCR (9). The resulting probes were labeled by nick-translation using biotin-16–dUTP (Boehringer Mannheim). After *in situ* hybridization to human metaphase chromosomes fixed by methanol/acetic acid, signals were detected via avidin-conjugated fluorescein isothiocyanate (Vector Laboratories, Burlingame, CA). For identification and band assignment, chromosomes were counterstained by DAPI (4,6-diamidine-2-phenylindole dihydrochloride). Through the use of a cooled CCD camera (Photometrix, Tucson, AZ), digitized images were generated for each of the two fluorochromes and overlaid electronically after digital processing. This allowed the assignment of the RanBP2 gene to chromosome 2q11–q13 (Fig. 1).

To obtain evidence for the presence of introns in the gene, PCR was performed using DNA from a panel of hamster somatic cell hybrids (Coriell Institute for Medical Research, Camden, NJ) under the following conditions: 30 cycles of amplification, 1 min of denaturation at 96°C, 30 s of annealing at 55°C, and 1 min of extension at 72°C. Each reaction was carried out in 50 μ l containing 10 mM Tris–HCl (pH 8.3), 2 mM MgCl₂, 50 mM KCl, 200 μ M each dNTP, 0.25 mM each of two primers, 50 ng of DNA template, and 1 U *Taq* DNA polymerase. Using the sense primer 5′-GCAGAATTGCTTTGTAA-3′ and the antisense primer 5′-AGT TCTGACTGAATCAAG-3′, we obtained a specific 4-kb product from both the YAC and a hybrid cell line harboring human chromosome 2, whereas no product was obtained from hybrids containing other human chromosomes. This result suggests the presence of a 3.75-kb intron in this region. The PCR product was sequenced, and an exon/intron transition was identified at position +405 of

the open reading frame (Genbank Accession No. D42063). Furthermore, we used an intron sequence primer, 5′-GAC-TAGTTGCTGAGGATTG-3′, to verify the presence of the intron. This confirms the localization of the gene for RanBP2 to chromosome 2q11–q13. Molecular genetic changes on 2q without identification of a specific gene have been described for human male germ cell tumors (5) and for neuroblastomas (8).

ACKNOWLEDGMENTS

We thank Sibylle Ohl for excellent technical assistance. This work was supported by the German–Israeli Cooperation in Cancer Research and by a grant from the Deutsche Forschungsgemeinschaft to H.P.

REFERENCES

1. Bischoff, F. R., Krebber, H., Smirnova, E., Dong, W. H., and Ponstingl, H. (1995). Co-activation of RanGTPase and inhibition of GTP dissociation by Ran GTP binding protein RanBP1. *EMBO J.* **14**: 705–715.
2. Church, G. M., and Gilbert, W. (1984). Genomic sequencing. *Proc. Natl. Acad. Sci. USA* **81**: 1991–1995.
3. Ferreira, P. A., Nakayama, T. A., Pak, W. L., and Travis, G. H. (1996). Cyclophilin-related protein RanBP2 acts as chaperone for red/green opsin. *Nature* **383**: 637–640.
4. Lichter, P., Tang, C. C., Call, K., Hermanson, G., Evans, G. A., Housman, D., and Ward, D. C. (1990). High resolution mapping of human chromosome 11 by *in situ* hybridization with cosmid clones. *Science* **247**: 64–69.
5. Lothe, R. A., Peltomaki, P., Tommerup, N., Fossa, S. D., Stenwig, A. E., Borresen, A. L., and Nesland, J. M. (1995). Molecular genetic changes in human male germ cell tumors. *Lab. Invest.* **73**: 606–614.
6. Mahajan, R., Delphin, C., Guan, T., Gerace, L., and Melchior, F. (1997). A small ubiquitin-related polypeptide involved in targeting RanGAP1 to nuclear pore complex protein RanBP2. *Cell* **88**: 97–107.
7. Sazer, S. (1996). The search for the primary function of the Ran GTPase continues. *Trends Cell Biol.* **6**: 81–85.
8. Takita, J., Hayashi, Y., Kohno, T., Shiseki, M., Yamagushi, N., Hanada, R., Yamamoto, K., and Yokota, J. (1995). Allelotype of neuroblastoma. *Oncogene* **11**: 1829–1834.
9. Wilgenbus, K. K., Mincheva, A., Korn, B., Lichter, P., and Poustka, A. (1995). IRS-long range (LR) PCR: A simple method for efficient amplification of human genomic DNA from complex sources. *Methods Mol. Cell. Biol.* **5**: 214–221.
10. Wilken, N., Senecal, J. L., Scheer, U., and Dabauvalle, M. C. (1995). Localization of the Ran-GTP binding protein RanBP2 at the cytoplasmic side of the nuclear pore complex. *Eur. J. Cell Biol.* **68**: 211–219.
11. Wu, J., Matunis, M. J., Kraemer, D., Blobel, G., and Coutavas, E. (1995). Nup358, a cytoplasmically exposed nucleoporin with peptide repeats, Ran-GTP binding sites, zinc fingers, a cyclophilin A homologous domain, and a leucine-rich region. *J. Biol. Chem.* **270**: 14209–14213.
12. Yokoyama, N., Hayashi, N., Seki, T., Panté, N., Ohba, T., Nishii, K., Kuma, K., Hayashida, T., Miyata, T., Aebi, U., Fukui, M., and Nishimoto, T. (1995). A giant nucleopore protein that binds Ran/TC4. *Nature* **376**: 184–188.