Centromere activity in dicentric small supernumerary marker chromosomes

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Abstract Twenty-five dicentric small supernumerary marker chromosomes (sSMC) derived from #13/21, #14, #15, #18, and #22 were studied by immunohis-tochemistry for their centromeric activity. Centromere protein (CENP)-B was applied as marker for all centromeres and CENP-C to label the active ones. Three different 'predominant' activation patterns could be observed, i.e., centric fusion or either only one or all two centromeres were active. In one inherited case, the same activation pattern was found in mother and son. In acrocentric-derived sSMC, all three activation patterns could be present. In contrary, in chromosome 18-derived sSMC, only the fusion

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T. Liehr (⊠) Institut für Humangenetik, Postfach, 07740 Jena, Germany e-mail: i8lith@mti.uni-jena.de type was observed. In concordance with previous studies a certain centromeric plasticity was observed in up to 13% of the cells of an individual case. Surprisingly, the obtained data suggests a possible influence of the sSMC carrier's gender on the implementation of the predominant activation pattern; especially, only one active centromere was found more frequently in female than in male carriers. Also, it might be suggested that dicentric sSMC with one active centromere could be less stable than such with two active ones-centromeric plasticity might have an influence here, as well. Also, centromere activity in acrocentric-derived dicentrics could be influenced by heteromorphisms of the corresponding short arms. Finally, evidence is provided that the closer the centromeres of a dicentric are and if they are not fused, the more likely it was that both of them became active. In concordance and refinement with previous studies, a distance of 1.4 Mb up to about 13 Mb the two active centromere state was favored. while centromeric distance of over ~15 Mb lead to inactivation of one centromere. Overall, here, the first and largest ever undertaken study in dicentric sSMC is presented, providing evidence that the centromeric activation pattern is, and parental origin may be of interest for their biology. Influence of mechanisms similar or identical to meiotic imprinting in the centromeric regions of human chromosomes might be present. Furthermore, centromeric activation pattern could be at least in parts meaningful for the clinical outcome of dicentric sSMC, as sSMC stability and

mosaicism can make the difference between clinically normal and abnormal phenotypes.

Keywords dicentrics \cdot small supernumerary marker chromosomes (sSMC) \cdot CENP-B \cdot CENP-C \cdot centromere activity

Abbreviations

ANOVA	Analysis of variance
CENP	Centromere protein
MB	Megabasepair
sSMC	Small supernumerary marker chromosome

Introduction

For polar movements of chromosomes during mitosis interaction of microtubules with a special centromere structure, the kinetochore, are necessary. A normal human chromosome harbors only one active centromere and derivative chromosomes with two or more centromeres (dicentrics or multicentrics) are generally thought to have only one active centromere during cell division. It is assumed that several active centromeres on one piece of DNA should lead to faulty alignments; two centromeres on one chromatid could orient to opposite poles, which would result in anaphase bridges and tearing of the chromosome. However, when the two centromeres are close together, there is little room for torsion between them, and stable dicentrics can be formed (Niebuhr 1972; Daniel and Lam-Po-Tang 1976; Dewald et al. 1979; Ing and Smith 1983; Rivera et al. 1989). Also, it was already shown that the presence of two functional kinetochores on a single chromosome does not invariably lead to chromosome instability and loss (Sullivan and Willard 1998). During human meiosis, rarely but regularly, a U-type exchange between different (Liehr et al. 2004) or within the same chromosome (Murmann et al. 2009; Sheth et al. 2009 take place. This leads to stable dicentric, mostly acrocentric-derived derivative chromosomes, so called dicentric small supernumerary marker chromosome (sSMC). At present, about 2.7 million carriers of such sSMC are alive, about two-thirds of which harbor dicentric, often mentioned as inverted duplicated (inv dup) sSMC (Liehr and Weise 2007).

Anticentromeric antibodies were identified in the sera of patients with the calcinosis, Raynaud's

syndrome, esophageal dysmotility, sclerodactyly, and telangiectasia (CREST) variety of scleroderma (Moroi et al. 1980). These sera recognize both centromeres in normal and dicentric chromosomes except for the Y chromosome (Merry et al. 1985; Earnshaw and Migeon 1985; Peretti et al. 1986; Rivera et al. 1989; Wandall 1989; Haaf and Schmid 1990). Different proteins were recognized by the CREST sera and their location in the centromere determined: centromere protein (CENP)-A is a centromere-specific histone similar to H3 (Palmer et al. 1991), CENP-B is distributed in the centromere region beneath the kinetochores (Cooke et al. 1990) where it binds to a recognition sequence in human alpha-satellite DNA (Masumoto et al. 1989), CENP-C is a component of the inner kinetochore plate (Saitoh et al. 1992), and CENP-D is similar to the RCC1 protein, a negative regulator of mitosis (Bischoff et al. 1990). However, only CENP-C differentiates between active and inactive centromeres (Earnshaw et al. 1989).

Here, we studied for the centromere activity in 25 dicentric sSMC.

Material and methods

Studied cases

Carnoy-fixed cell suspensions derived from peripheral blood of 25 carriers with a dicentric sSMC were included in the present study. As summarized in Fig. 1 and Table 1, three sSMC each were derived from chromosome 13 or 21 and chromosome 22, four each from chromosome 14 and 18, and the remainder from chromosome 15. The female-to-male ratio was 13 to 12. In 11 cases, the sSMC were de novo, in five maternally derived, in one case familial (maternal and paternally transmitted), and in the rest of the cases the origin was not known. An sSMC was present in 20-100% of the studied peripheral blood cells. The clinical data of the studied cases is available at (Liehr 2010).

Molecular cytogenetics

The origin and genetic content of each sSMC was determined by fluorescence in situ hybridization (FISH) as reported previously, using different multicolor-FISH approaches. Centromere-specific



Fig. 1 Chromosomal distribution of the studied 25 sSMC cases. *chr* chromosome

FISH in its different variants was applied (cenM-FISH: Liehr et al. 2006; acrocenM-FISH: Trifonov et al. 2003; subcenM-FISH: Liehr et al. 2006). Additionally, in inv dup (15) cases, the commercially available probe LSI UBE3A (Abbott) located in 15q13 was used according to manufacturer's instructions.

Immunohistochemistry

The immunoistochemical tests were done on Carnoyfixed cell suspension as previously published (Earnshaw et al. 1989). A rabbit polyclonal to CENP-B (Abcam, Cambridge, UK) was used to stain all centromeres (dilution 1:50). The specific staining of the active centromeres was performed with the anti-CENP-C antibody guinea pig serum (1:100; Ando et al. 2002). FITC-labeled goat anti-rabbit IgG and CyTM3-conjugated AffiniPure Goat Anti-Guinea Pig IgG (Dianova, Hamburg, Germany) were applied as secondary antibodies.

Chromosome banding was achieved by DAPI counterstaining (4'6-diaminidino-2-phenylindoleb, Sigma). Per case between 50 and 200 metaphases with sSMC were evaluated. An Axioplan 2 fluorescence microscope (Zeiss, Jena, Germany), a standard CCD camera (IMAC) and the software ISIS (Metasystems, Altlussheim, Germany) were used for the analysis.

Statistics

Statistical analysis was performed using Student's *t* test and one-way analysis of variance (ANOVA) method. Statistical significance was defined as p < 0.05.

Results

As summarized in Table 1 and Fig. 2 in the 25 studied cases, three different activation pattern of the two centromeres of each sSMC were found: in six cases, two centromeres were active ($2 \times$ active); in nine cases, only one of the two centromeres was active ($1 \times$ active); and in ten cases, the centromeres were fused to one active unit (fusion); for examples of the three patterns, see Fig. 3.

Results of correlations of the three detected activation pattern fusion, $1 \times$ active and $2 \times$ active with different parameters are shown in Fig. 4. No statistically significant differences could be found when comparing de novo and parentally inherited sSMC (Fig. 4a); after Student's *t* test all *P* values were between 0.101 and 0.599 (detailed results not shown).

However, gender-dependent activation patterns were found: the states fusion and $2\times$ active are found more frequently in male, while $1\times$ active is present in seven female compared to only two male (Fig. 4a). This is a statistically significant difference with a *P* value of 0.012 according to Student's *t* test. Fusion and $2\times$ active *P* values were 0.372 and 0.247.

Statistically significant differences were also found for sSMC mosaicism and centromere activity (Fig. 4c). A fusion or two active centromeres (2× active) was present mostly in non-mosaic: Student's *t* test *P* values were 0.003 (fusion), 0.011 (2× active), and <0.001 (1× active). This means sSMC with 2× active were less stable, while in almost all of the cases with only one active centromere (1× active) the sSMC was stable in >90% of the peripheral blood cells. This was supported by ANOVA test showing a statistically significant difference for mosaic and non-mosaic cases: F=10.744, P=<0.001.

Finally, there was a possible correlation of the distance of both centromeres and their activity. The closer the centromeres were located, the more likely either they were fused or two of them were active (see Fig. 4d).

Discussion

The present study is the first systematic study performed in not less than 25 dicentric sSMC. Only one comparable larger study was done in 15 cases with Robertsonian translocations (Page and Shaffer

Table 1 Summary of gender data with immunohistochemical, cytogenetic, and molecular data of studied sSMC cases

Cases numbered according to Liehr 2010	Gender	Parental origin	Cells with sSMC (%)	sSMC characterized as	Centromere distance (MB)	One centromere inactive in (%) of cells with sSMC	Interpretation of immuno- histochemistry results
13/21-U-22	Female	n.a.	100	inv dup (13 or 21)	n.a.	n.a.	Fusion
13/21-O-q10/1-11	Male	dn	100	(q_{10}) inv dup (13 or 21) (q_{10})	n.a.	n.a.	Fusion
13/21-O-q11/2-5	Male	n.a.	20	(q10) inv dup(13)(q11) or inv dup (21)	n.a.	21	2× active
14-O-q11.1/1-6	Male	n.a.	20	$(q_{11,1})$ inv dup(14) $(q_{11,1})$	n.a.	n.a.	Fusion
14-CO-5	Female	mat	57	$(q_{11,1})$ inv dup(14) $(q_{11,1})$	n.a.	97	$1 \times$ active
14-O-q11.1/1-10	Female	n.a.	52	(q11.1) inv dup(14) (q11.1)	n.a.	91	$1 \times$ active
14-O-q11.1/1-22	Male	dn	96	(q11.1) inv dup(14) (q11.1)	n.a.	8	2× active
15-O-q11.1/1-63	Female	fam	83	(q11.1) inv dup(15)	n.a.	98	$1 \times$ active
15-U/8-4	Female	dn	56	(q11.1) inv dup(15)	n.a.	97	$1 \times$ active
O15-q11.1/1-8	Female	dn	99	(q11.1) inv dup(15) (z11.1)	n.a.	13	2× active
15-O-q11.1/1-11	Male	mat	98	$(q_{11,1})$ inv dup(15)	n.a.	2	2× active
15-O-q11.1/1-16	Female	n.a.	28	(q11.1) inv dup(15) (q11.1)	n.a.	100	$1 \times$ active
Son of				(q11.1)			
15-O-q11.1/1-16	Male	mat	73	inv dup (15)	n.a.	91	1× active
15-O-q11.1/1-45	Female	n.a.	52	$(q_{11,1})$ inv dup(15) $(q_{11,1})$	n.a.	n.a.	Fusion
15-O-q11.2~12/1-	Female	n.a.	87	$(q_{11,1})$ inv dup(15) $(q_{11,2},12)$	~4.5	2	2× active
15-CWw-148	Male	dn	55	$(q_{11,2}, q_{12})$ inv dup(15) (q_{12}, q_{13})	15.4	98	1× active
15-W-q13/2-4	Female	dn	23	(q_{12}, q_{13}) inv dup(15)(q13) x ²	19.5	100	1× active
15-W-q14/1-6	Female	dn	99	inv dup(15)(q14)	~27	100	1× active
18-Wi-132	Male	dn	100	inv dup (18)	n.a.	n.a.	Fusion
18-Wi-143	Male	dn	100	$(q_{11,1})$ inv dup(18) $(q_{11,1})$	n.a.	n.a.	Fusion
18-Wi-142	Female	dn	100	$(q_{11,1})$ inv dup(18) $(q_{11,1})$	n.a.	n.a.	Fusion
18-Wi-146	Female	dn	100	$(q_{11,1})$ inv dup(18) $(q_{11,1})$	n.a.	n.a.	Fusion
22-O-q11.1/1-5	Male	mat	100	$(q_{11,1})$ inv dup(22) $(q_{11,1})$	n.a.	n.a.	Fusion
22-U-41	Male	mat	100	$(q_{11,1})$ inv dup(22) $(q_{11,1})$	n.a.	n.a.	Fusion
22-Wces-5-95	Male	dn	100	(q11.1) inv dup(22) (q11.21)	1.4	4	2× active



Fig. 2 Three different activation pattern as detected in the 25 studied cases: fusion, one centromere active (1× active) and both centromeres active ($2 \times$ active)

1998) using also CENP-B as all-centromere marker and CENP-C for active centromeres (Earnshaw et al. 1989). CENP-C is either necessary for anaphase chromosome movement or for mediating a signal which triggers centromere function during anaphase (Fukagawa and Brown 1997), however, it is necessary but not sufficient for the formation of a functional centromere (Fukagawa et al. 1999).

In the studied 25 dicentric sSMC, three different activation patterns could be present (fusion, $1 \times$ active, $2\times$ active). In concordance with previous studies (Wandall 1994; Page and Shaffer 1998; Higgins et al. 2005), a certain centromeric plasticity could be observed; i.e., in between 0% and 13% of the cells of an individual case the sSMC showed another centromere-activity pattern than the majority, which is called in the following 'predominant pattern' (see Table 1, second to last column). Notable, centromeric plasticity was not observed if a fusion was present. Case 15-O-q11.1/1-16 and son of case 15-O-q11.1/1-16 provide first evidence that in familial cases an identical activation pattern could be retained. This is especially remarkable as in the mother (case 15-Oq11.1/1-16) the cells with an sSMC were by far less frequent in the peripheral blood than in the son. However, it is known that sSMC can be lost in peripheral blood throughout live time (Liehr et al. 2004). As visible in Table 1, all four inv dup(18) (q11.1) cases showed functional centromeric fusion, while in the acrocentric-derived sSMC in principle all three activation patterns could be present.

Here, for the first time, evidence was provided that the predominant centromere activation pattern is dependant from the gender of the sSMC carrier: fusion and 2× active are found more frequently in male, while $1 \times$ active is more frequent in female. This still could be due to an ascertainment bias and low the



Fig. 3 Examples for the three different activation pattern: both centromeres active (2× active) were found in case 15-O-q11.1/ 1-22-CENP-B (green labeling all centromeres) and CENP-C (red labeling only active centromeres) label simultaneously the centromeres on normal chromosomes 15 (#15) and both



centromeres of the dicentric sSMC. In case 15-O-q11.1/1-63 and 22-O-q11.1/1-5, only the sSMC are shown; the first shows only one active centromere $(1 \times \text{ active})$, the latter a typical fusion of green and red signal





Fig. 4 a-c Comparison of the three observed three predominant immunohistochemical pattern fusion, $1 \times$ active and $2 \times$ active with inheritance (inherited or de novo, **a**), gender (male or

female, **b**) and sSMC stability (mosaic or non-mosaic, **c**). **d** Alignment of chromosome size (expressed in chromosomal band) and only one active centromere

number of examined cases; further studies are necessary to verify that suggestion, even though the difference was statistically significant. If true, this observation might be helpful to explain the fact that familial sSMC are transmitted predominantly via the maternal line (Liehr 2006). It is considered that a dicentric chromosome with only one active centromere is more stable than one with two active centromeres (Therman et al. 1986). If the latter would be applicable also to dicentric sSMC during meiosis, the fact that dicentric sSMC with one active centromere are more frequently present in female could be of interest. Controversy, the present data show an adverse influence of $1 \times$ active at least on the mitotic sSMC stability (Fig. 4c). Additionally, four of the six sSMC with two active centromeres were mitotically stable (no mosaic formation) in the present study which might also be due in parts to the here and previously observed centromeric plasticity (Wandall 1994; Page and Shaffer 1998; Higgins et al. 2005).

A study in larger dicentric X chromosomes showed that the centromere activity is correlated to the centromere distance, i.e., the closer the centromeres, the more likely it was that both of them became active (between 4 and 12 Mb), while in two distant centromeres, only one was active (34 MB; Sullivan and Willard 1998). Here, in principle, this trend was confirmed (see Table 1 and Fig. 4d), even though reliable data on the centromere distance was available only for five of the cases. If centromeric distance was over 15.4 Mb, only one centromere was active (cases 15-CWw-148, 15-W-q13/2-4 and 15-W-q14/1-6). In those cases where the centromeric distance was between 1.4 and ~4.5 Mb (22-Wces-5-95 and 15-Oq11.2~12/1-2), two active centromeres could be observed. In the remainder 20 cases, the sSMC consisted according to molecular cytogenetics in their q-arms only of heterochromatic material and were broken in band q10 to q11.1. Both cases with cytogenetic breakpoint in q10 showed fusion pattern of centromere activity; besides, also eight cases with breaks in q11.1 had fusion pattern. The remainder ten q11.1 cases showed two clearly separated centromeric signals using the CENP-B antibody; four of those showing predominantly the 2× active pattern applying CENP-C antibody. Here, centromere plasticity may be not enough to explain this variance. However, it is a well-known fact that the acrocentric short arms are highly heteromorphic. Rearrangements in the short arms like reported previously (Lau et al. 1979; Schmid et al. 1994; Friedrich et al. 1996; Verma et al. 1996; Reddy and Sulcova 1998) may be causative for the observed variance. Thus, this might explain also, that cases with the same karyotype, like 47,XN,+inv dup(15)(q11.1) could have any of the three pattern fusion (case 15-O-q11.1/1-45), $1 \times$ active (case 15-O-q11.1/1-16) and $2 \times$ active (case 15-O-q11.1/1-11).

Conclusion

By this first and largest ever undertaken study in dicentric sSMC, evidence is provided that the centromeric activation pattern is of interest for the biology of sSMC maintenance and stability in individuals. Influence of mechanisms similar or identical to meiotic imprinting could act upon also the centromeric regions of human chromosomes, as possibly visible in the male-to-female ration in dicentric sSMC cases with one active centromere only. Furthermore, interesting clinical impact of mosaicism in sSMC carriers was reported: sSMC with known adverse prognosis if present in 100% of the cells of a carrier were observed to be harmless if present in small mosaic state only (Bonati et al. 2005; Loitzsch and Bartsch 2006; Guichet et al. 2009). Thus, understanding of centromeric activity and possible influence on sSMC stability have to be studied further also for the establishment of proper genotype-phenotype correlations in sSMC.

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