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Serum levels of S-100 protein are directly proportional to the size, number, thickness and degree of cellularity of congenital melanocytic nevi

To the Editor: Some patients with congenital melanocytic nevi (CMN) present progressive growth and thickening, extracutaneous involvement (neurocutaneous melanocytosis, NCM) or neoplastic transformation (melanoma); and others remain stable or even regress. There are no markers to assess progression or follow-up. Recently, we found S-100, a protein which acts on cell differentiation and proliferation, elevated in CMN.¹ S-100 is a ligand of the RAGE pathway (related to the MAPK-pathway), and low serum levels of soluble-RAGE were related to poor survival in melanoma.² Also SOX10, expressed in melanocytes with high specificity, is useful in detection, prognosis and treatment assessment of melanoma.³ We explored if S-100, RAGE and SOX10 serum levels vary in children's CMN and assessed clinical or pathological correlations.

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Patients (N = 64)		S-100 (μg/L): mean (IQR)	P value
Age, median (IQR)	9 (5-12)		
Male, n (%)	21 (32.81)		
Neurocutaneous melanosis, n (%)			.843
No	56 (87.5)	0.31 (0.13-0.55)	
Yes	8 (12.5)	0.27 (0.14-0.46)	
Subcutaneous tissue alteration, n (%)			.042
No	55 (85.9)	0.19 (0.10-0.38)	
Yes	9 (14.1)	0.35 (0.19-0.71)	
Size classification depending on PAS n (%)			.030
M1 1.5-10 cm	11 (17.19)	0.09 (0.05-0.14)	
M2 >10-20 cm	8 (12.50)	0.19 (0.14-0.45)	
L1 >20-30 cm	11 (17.19)	0.31 (0.16-0.58)	
L2 >30-40 cm	5 (7.81)	0.31 (0.11-0.71)	
G1 >40-60 cm	12 (18.75)	0.17 (0.08-0.51)	
G2 >60 cm	17 (26.56)	0.31 (0.19-0.43)	
No. of satellite nevi n (%)			.039
No satellites	23 (35.94)	0.14 (0.09-0.25)	
<20 satellites	14 (21.88)	0.28 (0.12-0.52)	
20-50 satellites	11 (17.19)	0.49 (0.15-1.35)	
>50 satellites	16 (25.00)	0.31 (0.19-0.46)	
Dermal or subcutaneous nodules <i>n</i> (%)			.087
None	41 (64.06)	0.19 (0.10-0.32)	
Scattered	14 (21.88)	0.27 (0.14-0.52)	
Extensive	9 (14.06)	0.53 (0.19-1.21)	
Hypertrichosis n (%)			.043
None	8 (12.50)	0.09 (0.05-0.19)	
Notable	25 (39.06)	0.27 (0.11-0.75)	
Marked	31 (48.44)	0.22 (0.15-0.49)	
General location #n (%)			.588
Head	25 (39.06)	0.16 (0.09-0.32)	
Trunk	10 (15.63)	0.26 (0.09-0.76)	
Extremities	7 (10.94)	0.15 (0.13-0.78)	
Head $+$ trunk	4 (6.25)	0.22 (0.09-1.08)	
Trunk + extremities	16 (25.00)	0.32 (0.16-0.50)	
Head $+$ trunk $+$ extremities	2 (3.13)	0.38 (0.22-0.53)	

Table I. Characteristics of CMN patients and lesions and clinical and histopathological correlation with serum protein levels

Histopathological analysis (N = 25)	Melanocyte extent	Serum protein	Rho	P value
Deep dermis	Melan-A	S-100	0.41	.043
		SOX10	0.64	<.001
		RAGE	0.18	.384
	S-100	S-100	0.44	.028
		SOX10	0.58	.002
		RAGE	-0.17	.421
Subcutaneous tissue	Melan-A	S-100	0.34	.107
		SOX10	0.29	.170
		RAGE	-0.20	.334
	S-100	S-100	0.61	.001
		SOX10	0.38	.080
		RAGE	-0.06	.765

Bold P-values indicate statistically significant results; bold S-100 levels and IQR in clinical features indicate the subgroups significantly different.

Regarding histopathological analysis, Melan-A⁺ and S-100⁺ areas and the number of Ki-67⁺ cells were studied using ImageJ software in dermis and with conventional semi-quantitative scales in subcutaneous tissue. In addition to results presented in the table, a positive correlation between serum SOX10 and subcutaneous Ki67⁺ (Rho 0.438 P = .029) and a negative correlation between serum RAGE and lesion thickness (Rho -0.62 P = .001) were found. Supplementary Methods Material, available via Mendeley at https://data.mendeley.com/datasets/mzr3rvxdkh/1. *CMN*, Congenital melanocytic nevus; *G1 and G2*, giant; *IQR*, interquartile range; *L1 and L2*, large; *M1 and M2*, medium; *PAS*, projected adult size.



Fig 1. Serum determinations of S-100, RAGE and SOX10. HC: healthy children (27 men [42.18%] P = .215 vs CMN; median age: 9 [IQR 6-12] P = .464 vs CMN). MM: metastatic melanoma patients with disease progression (4 men [40%]; median age: 62.5 [IQR 57-74]). Healthy controls were children seen at the emergency department who underwent serologic testing for suspected infections, and metastatic melanoma patients were undergoing laboratory review. A, Comparison of levels of S-100 between HC, CMN and MM. CMN patients showed higher levels of S-100 (mean 0.38; standard deviation [SD] 0.43 ug/L; mean for HC 0.08 [SD 0.05] ug/L; and mean for MM 0.56 [SD 0.88] ug/L; P < .001). **B**, Comparison of levels of **RAGE** between HC, CMN and MM. CMN patients showed lower levels than HC and higher than MM (mean for CMN 1805.10 [SD] 1098.89 pg/mL; mean for HC 2189.07 [SD 857.20] pg/mL; and mean for MM 1337.85 [SD 634.71] pg/mL; P < .001). C, Comparison of levels of SOX10 between HC, CMN and MM. SOX10 was higher in CMN than in HC patients (mean for CMN 1316.84 [SD] 768.49 pg/mL; mean for (HC) 895.21 [SD 222.39] pg/mL; P < .001). When comparing SOX10 between CMN and MM (mean for MM 858.73 [SD] 379.54 pg/mL) no significant differences were found (P = .200). Some of the CMN patients had already undergone surgeries when they entered the study. The percentage (median, IQR) of the CMN present/maintained when the blood samples were collected was 85, 50-100. The most frequent surgeries performed when obtaining samples were expanders and tissue removal + flap (36%), tissue excision (32%) and expanders placement (19.5%). Samples were taken before starting the procedure.

This prospective, case-control, multicenter study enrolled consecutive patients with mediumto-giant-CMNs who underwent surgery (April 2019-December 2020). Left over serum samples of sex-and-age-matched healthy controls (HC) and a small group of metastatic melanoma patients were collected. The study was approved by the University Clinic of Navarra institutional review board and participants provided informed consent. CMN categorization was performed according to Krengel et al⁴ criteria. S-100 was determined with an electrochemiluminescence-immunoassay; RAGE and SOX10 with ELISA kits. Representative tissue specimens taken during certain patients' surgery were analyzed for nevomelanocytic extent and proliferation. Supplementary Material, available via Mendeley at https://data.mendeley.com/ datasets/mzr3rvxdkh/1.

64 CMN (Table 1), 64 HC and 10 metastatic melanoma patients were recruited. CMN presented the highest levels of S-100; and lower levels of RAGE and higher levels of SOX10 than HC (Fig 1). Serial analyses were performed in 11 CMN showing that S-100 and SOX10 tended to decrease (correlation S-100-SOX10 Rho 0.253, P = .362) while RAGE increased (correlation S-100-RAGE Rho -0.432, P = .107) across

time and treatments (months between determinations: median 21, 7-36). S-100 showed association with size, satellite nevi, hypertrichosis and subcutaneous tissue alterations (Table I and Supplementary Fig 2, available via Mendeley at https://data.mendeley.com/datasets/ mzr3rvxdkh/1), and multivariate analysis showed that extensive dermal nodules, >20 satellites, or at least notable hypertrichosis increased S-100 values (coefficients 0.35, P = .011; .34, P = .008; and .20, P = .041). Higher serum levels of S-100 and SOX10 correlated with higher Melan-A⁺ and S-100⁺deep dermis areas; and S-100 serum levels correlated with S-100⁺ subcutaneous-areas (Table I). Serum SOX10 and subcutaneous Ki67⁺ presented positive correlation (Rho 0.438 P = .029) and serum RAGE presented negative correlation with lesion thickness (Rho -0.62P = .001).

Findings indicate higher nevomelanocytes populations result in greater S-100 release into the blood, as patients with larger, multiple, deeper and more cellular lesions presented higher levels. CMNs' size and satellites were reported to correlate with melanoma and NCM though no definitive conclusions could be made in regard to these complications in the present study due to sample size. However, it seems plausible that in those children affected by CMN suffering overgrowth, melanoma, or NCM progression, serial determinations of serum S-100 could help in early detection and treatment, being complementary to physical examination and MRI. Usefulness of serum S-100 in melanoma, even within normal ranges, was highlighted recently.⁵

Long-term studies are needed to understand serum proteins variations across time in different settings. S-100 could be an easy, noninvasive, affordable analysis in CMN characterization and follow-up.

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Conflicts of interest

None disclosed.

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Time spent outdoors and sun protective behaviors among sexual minority women

To the Editor: Sexual minority women (SMW; eg, lesbian, bisexual), a National Institutes of Health US health disparity population,¹ may have a lower prevalence of indoor tanning and skin cancers compared to heterosexual women^{2,3}; however, the relationship between sun protective behaviors (SPBs), sexual orientation, and sexual behavior is not well-studied. This cross-sectional study compared SPBs in women by sexual orientation and sexual behavior using nationally representative National Health and Nutrition Examination Survey data pooled from 2009 to 2016.

Self-reported sexual orientation was dichotomized into SMW (lesbian, bisexual) and heterosexual; sexual behavior was dichotomized into