

RAS pathway influences the number of melanocytic nevi in cardiofaciocutaneous and Costello syndromes

To the Editor: The number of melanocytic nevi is one of the strongest risk factors for melanoma,¹ yet the reasons for the interpersonal variability in this number are largely unknown. Both nevi and melanoma show somatic mutations in the RAS pathway components, most commonly in *BRAF*.² Germline mutations in these same genes cause a group of developmental syndromes termed RASopathies,³ some of which characteristically display melanocytic nevi.⁴ However, the effects of germline RAS pathway mutations on the number of nevi are poorly understood.

To determine how the number of melanocytic nevi is influenced by the RAS pathway, we analyzed the numbers of nevi in cardiofaciocutaneous syndrome (CFC) and Costello syndrome (CS). CFC is caused by mutations in the downstream elements of the RAS pathway, including *BRAF*, *MAP2K1*, and *MAP2K2*, or, rarely, *KRAS*,^{5,6} whereas CS results from mutations in an upstream core component of the pathway, *HRAS* (Fig 1).⁷

The institutional review board at the University of California Davis approved the study. After informed

consent, 16 individuals with CFC and 24 with CS were enrolled.⁸ Photography of nevi was performed (Canfield Scientific, Parsippany, NJ). The photographs were examined for the number of nevi by 2 authors independently and without knowledge of the mutation status. Fitzpatrick skin phototype was determined based on the photographs and a questionnaire regarding tendency to burn and tan. A 2-sample *t* test and linear regression models were used.

The mean age for individuals with CFC was 15.1 years (range, 6-35 years) and for those with CS was 14.5 years (range, 6-31 years).⁸ The majority of individuals with CFC reported a *BRAF* mutation (*BRAF* in 13/16 [81.3%], *MAP2K1* in 1/16 [6.3%], *MAP2K2* in 1/16 [6.3%], and unknown in 2/16 [12.5%]), and the majority of individuals with CS reported an *HRAS* mutation (p.G12S in 16/25 [64.0%]; p.G12C in 2/25 [8.0%]; p.G12A, p.G13C, p.G13D, p.A146V, and p.K117R in 1 each/25 [4.0%]; and unknown in 2/25 [6.3%]).⁸

A marked difference was noted in the number of nevi in CFC versus CS (Fig 2 and Table 1). The average number of nevi on the back was 47.8 in CFC (standard error of the mean [SEM], 14.0) and 8.1 in CS (SEM, 1.8; *P* = .002). The number of nevi in CS corresponds to published population-based data of 8.4 nevi on average on the back of children and

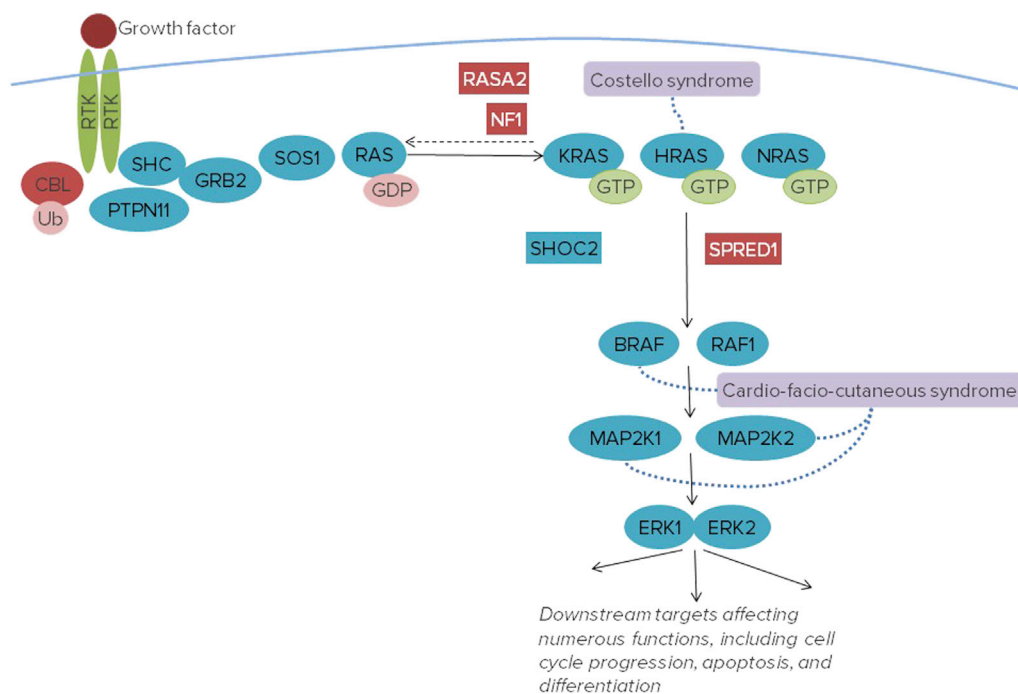


Fig 1. The RAS pathway showing genes mutated in patients with cardiofaciocutaneous syndrome and Costello syndrome in this study.



Fig 2. Increase in the number of nevi in (A) an individual with cardiofaciocutaneous syndrome compared with (B) an individual with Costello syndrome. Both individuals are women ages 22 to 23 years.

Table I. Number of nevi in CFC and CS

Characteristics	CFC	SEM or range	CS	SEM or range
Age, y	15.1	6-35	14.5	6-31
Number of nevi on back	47.8	14	8.1	1.8
Phototype I	6.0	n/a*	n/a	n/a
Phototype II	77.0	17.3	2.4	0.4
Phototype III	53.7	31.5	9.2	2.6
Phototype IV	16.0	5.0	10.3	3.7
Phototype V	15.0	n/a*	35.0	n/a*
Phototype VI	4.0	n/a*	9.0	n/a*
Number of nevi on face	24.3	7.3	4.0	0.8

CFC, Cardiofaciocutaneous syndrome; CS, Costello syndrome; n/a, not applicable; SEM, standard error of the mean.

*Only 1 individual; therefore, SEM is not applicable.

adolescents in the United States.⁹ The average number of nevi on the face was also increased in CFC (CFC: 24.3 [SEM, 7.3]; CS: 4.0 [SEM, 0.8]; $P = .001$). The number of nevi was higher for patients with older age (beta estimate, 1.8; 95% confidence interval, 0.3-3.3; $P = .02$). Moreover, the number of nevi was significantly higher for phototypes I through III in CFC compared with CS but not for phototypes IV through VI ($P = .01$). There were no significant differences in painful sunburns ($P > .99$), sunbathing habits ($P > .99$), or hours spent outdoors between CFC and CS patients ($P = .48$).

BRAF p.V600E is a well-known somatic driver of nevogenesis. This study expands our knowledge of germline regulators of nevus count by suggesting that germline mutations in *BRAF*, *MAP2K1*, and *MAP2K2*, but not the upstream core component of the RAS pathway *HRAS*, predispose to and influence the number of nevi. Moreover, increased numbers of nevi, especially in individuals with phototypes I

through III in CFC, suggest that ultraviolet (UV) radiation may enhance the effects of the germline mutations in the downstream components of the RAS pathway on nevogenesis. The results increase our knowledge of the genetic background and development of nevi, the potential precursors of melanoma. Although future studies are warranted to determine whether the risk of melanoma is increased in CFC, protection from UV radiation and regular skin examinations are appropriate for individuals with CFC.

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