Vettes in the Vineyard 2018



Chaumette Vineyards and Winery Ste. Genevieve County, MO

Saturday May 19, 11:00am to 3:00pm

\$25 per person includes entry fee; food; glass of wine, bottled beer, or soft drink; and donation!

Special "Best of the Vineyard" award presentation!

- Proceeds and donations Supporting PRISMS*
- Sponsored by the St. Louis Corvette Club
- People's Choice

***RSVP to tnrluebcke@charter.net OR 573-218-9051 by May 1!

From St. Louis: Take I-55 S to Exit 150, turn right for .1m, then left (S) onto Hwy B for 14 miles to Coffman. Veer left onto Hwy WW for 2.6 miles, Chaumette is on your right.

From Cape Girardeau: Take I-55 N to Exit 150, turn left for .2m, then left (S) onto Hwy B for 14 miles to Coffman. Veer left onto Hwy WW for 2.6 miles, Chaumette is on your right.

From Farmington: Take Hwy OO, to left on Hwy F, for 13 miles to Coffman. Turn right onto Hwy WW for 2.6 miles, Chaumette is on your right

- * Turn over for details
- **NO outside food or drink, please

WHAT is PRISMS??

PRISMS is the acronym for Parents and Researchers interested in Smith-Magenis Syndrome (SMS)

Summary:

Smith-Magenis syndrome (SMS) is a complex developmental disorder that affects multiple organ systems of the body. The disorder is characterized by a pattern of abnormalities present at birth (congenital) as well as behavioral and cognitive problems.

Common symptoms include distinctive facial features, skeletal malformations, varying degrees of intellectual disability, speech and motor delays, sleep disturbances, and self-injurious or attention-seeking behaviors. The specific symptoms present in each case can vary dramatically from one individual to another.

Approximately 90% of cases are caused when a portion of a chromosome is missing or deleted (monosomic). This deleted portion within chromosome 17p11.2 includes the RAI1 gene, which is believed to play a major role in the development of the disorder. Other genes within the deleted segment may also play a role in variable features in the syndrome, but it is not fully understood how significant a role they play in the development of SMS. In the remaining cases, there is no deleted material on chromosome 17; these cases are caused by mutations in the RAI1 gene.