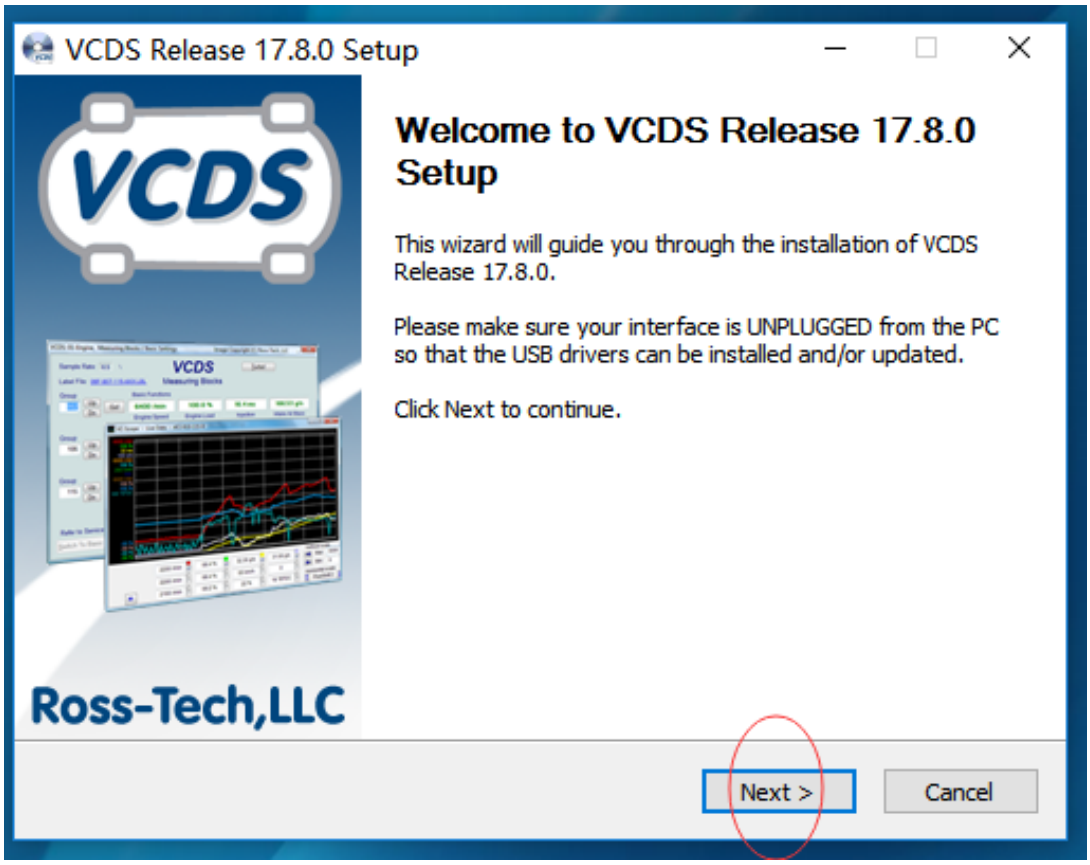
# Vag Com 12.10.3 Cracked Software Free Download

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. . "*A code was to be registered in the CIN10 database if the women's report to the service of a vaginal discharge was similar to a positive diagnosis of "condyloma acuminata" from any of the study physicians. During the study period, the participating physicians answered a detailed questionnaire about their clinical practice. In addition, reports of the VAG COM service were collected from the service provider database and entered into the database of the CIN10 project. * . . . "*In case a woman was lost to follow-up, either due to death or because the woman did not answer the invitation letter, a new invitation was sent after a 5-year interval. The observation period was terminated at the time of the last invitation to participate. The women answered questionnaires about lifestyle factors, reproductive history and experience, sexual behavior, and socio-demographic characteristics. * . . . "* The VAG COM study population was described according to demographic data, medical risk factors and conditions, family history of cervical cancer, reproductive factors, sexual behavior, and socio-economic status (SES). We used a set of questions to determine the woman's socioeconomic status, which was categorized into three groups: low, middle, or high SES. Middle SES was defined as a household income of less than two times the German minimum wage and high SES as a household income of more than twice the German minimum wage \[[@CR28]\]. With regard to family history, data were obtained on the prevalence of *p53* mutations and HPV type 16. A woman was considered to have a *p53* mutation when she had one or more cervical cancer-associated alterations in the *p53* gene (codons K100, K110, K117, T81, Y90, Y196, R175, R248, R273, and R282), *p53* gene amplification, homozygous *p53* gene deletion, or *p53* immunostaining in the biopsy material. A woman was considered to have HPV type 16 when she had the HPV 16 DNA sequence in the cervical sample. Laboratory techniques {#Sec4} ---------- We isolated the HPV-DNA from cervical scrapings with the High Pure PCR Template Preparation Kit (Roche Applied Science, Mannheim, Germany) using the instructions provided by the manufacturer. The Nested-PCR was carried out 82157476af