

# Sleep (Life Xpansion Code)

Client

**Demo Client** 

File

Module 1

# **Ideal Sleep Window**

The Ideal Sleep Window is the optimal period of time to get the most restful and rejuvenating sleep based on circadian rhythm. Circadian rhythm refers to the internal clock that regulates various biological processes in the body, including sleep-wake cycles. Every person has a unique circadian propensity, and it determines levels of alertness and restfulness throughout the day.

There are three categories of individuals associated with varying sleep windows: morning larks, intermediate, and night owls. Morning



larks are early risers who feel most alert during the first half of the day, with the earliest sleep window. On the other hand, night owls tend to feel more alert in the evening and night hours and have a later sleep window. Intermediate types fall somewhere in between morning larks and night owls.

When individuals regularly sleep outside of their ideal sleep window, performance can diminish significantly. Understanding one's ideal sleep window can provide valuable insights into how the body responds to different sleep patterns, and provide the opportunity to create the optimal sleep schedule.

# **Ideal Sleep Window**



### **Understanding what this may mean**

Intermediate sleep window means there's not a strong propensity towards being a night owl or an early morning lark. This is a relatively normal genetic result. In general, studies have found increased benefit from shifting to earlier sleep and wake times, but in general, people with this result tend to do best with bedtimes and wake-times of 9pm-12am and 5-8am.

# Genetic Summary 10

| SNP           | Marker      | Genotype | SNP          | Marker      | Genotype |
|---------------|-------------|----------|--------------|-------------|----------|
| CSNK1D T130C  | rs104894561 | TT       | CLOCK A*213G | rs1801260   | AG       |
| PER3 T1916G   | rs10462020  | TG       | CRY1 G636A   | rs8192440   | GG       |
| PER3 A3473G   | rs10462021  | AG       | PER3 C2566G  | rs228697    | CC       |
| PER2 T1984C   | rs121908635 | TT       | PER3 A1247G  | rs139315125 | AA       |
| CRY1 T1657+3G | rs184039278 | TT       | PER3 C1240G  | rs150812083 | CC       |

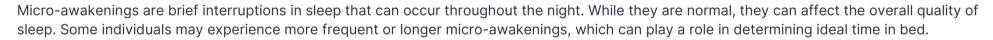
### Module 2

# **Time in Bed**

Getting the right amount of sleep is essential for overall health and well-being. There are many different aspects to time in bed; there's the time it takes to fall asleep, our actual sleep length, the micro-awakenings that occur each night, and are forgotten, and the true disruptions in sleep which wake us up.

Sleep length refers to the total amount of time spent sleeping, not including the time it takes to fall asleep or any awakenings during the night. Most people tend to focus on recommendations for sleep

length as recommendations for time in bed, which can result in consistently low sleep length and quality.



This page provides genetic propensities for the most ideal sleep length and time in bed, as well as the expected amount of micro-awakenings.



# **Ideal Total Time in Bed**

Neutral - 7.5 to 8 hours

### Understanding what this may mean

Individuals with this result may have a genetic propensity for requiring an average amount of time in bed. Maintaining good sleep hygiene, such as establishing a regular sleep schedule and creating a relaxing sleep environment, can help optimize the benefits of this amount of sleep.

# **Sleep Length Propensity**

# **Shorter Sleep**

### **Understanding what this may mean**

Individuals with this result may have a genetic propensity for requiring slightly less sleep than the average person. They may be able to function effectively on slightly less sleep than others, but it is important to note that chronic sleep deprivation can lead to negative health consequences. It is recommended that individuals with this result assess their sleep quality and ensure they are getting sufficient restorative sleep.

# **Micro-Awakenings Propensity**



### **Understanding what this may mean**

Individuals with this result may experience up to 75-90 minutes of micro-awakenings during the night, which may indicate a genetic propensity for more fragmented and less restful sleep. To optimize sleep, individuals with this result may benefit from practicing good sleep hygiene, such as establishing a consistent sleep schedule, creating a relaxing sleep environment, and addressing any underlying sleep disorders that may be contributing to the increased micro-awakenings.

# **Genetic Summary 9**

SNP Marker Genotype SNP Marker Genotype

| DEC2 G1151C       | rs121912617 | GG |
|-------------------|-------------|----|
| COMT G472A        | rs4680      | AA |
| CLOCK A*213G      | rs1801260   | AG |
| ABCC9 G3474-1222A | rs11046205  | GG |
| CLOCK A793-1130G  | rs11932595  | GG |

| CLOCK G-446-4252A   | rs12649507 | AG |
|---------------------|------------|----|
| BTBD9 G1154+104406A | rs3923809  | AG |
| MEIS1 T965+6302G    | rs2300478  | TT |
| PTPRD G103-113009A  | rs1975197  | AG |

### Module 3

# **Depression**

Depression, a complex and multifaceted mental health condition, is influenced by a myriad of factors, including genetic predispositions, environmental stressors, and lifestyle choices. This section delves into the genetic underpinnings of depression, aiming to shed light on how genetic variations can influence an individual's propensity for developing depression and guide personalized intervention strategies.



Research has identified several genetic markers associated with an increased risk of depression. Variations in genes related to neurotransmitter systems, such as serotonin and dopamine, play significant roles in mood regulation and can affect an individual's vulnerability to depression. Additionally, genes involved in the hypothalamic-pituitary-adrenal (HPA) axis, which regulates stress response, may contribute to the development and severity of depressive symptoms. The relationship between genetics and depression is further complicated by environmental influences, such as life stressors, trauma, and social support networks. Genetic predispositions can interact with these external factors, modulating the risk and expression of depression.

Identifying genetic predispositions to depression can inform personalized approaches to mental health care, including preventative measures, lifestyle modifications, and tailored treatment plans. For instance, individuals at higher genetic risk may benefit from early intervention strategies, such as stress management techniques, regular physical activity, and enhanced social support, to mitigate the impact of stressors and reduce the likelihood of developing depression. Genetic testing can also guide the selection of antidepressant medications and psychotherapy approaches, enhancing treatment efficacy and reducing trial-and-error processes. Pharmacogenomics, the study of how genes affect a person's response to drugs, is particularly promising in identifying which antidepressants are likely to be most effective or have fewer side effects for a given individual, based on their genetic makeup.

# **Depression Propensity**



### **Understanding what this may mean**

Individuals with a very low risk of depression have genetic markers that significantly reduce their likelihood of experiencing depression compared to the general population. This favorable genetic predisposition suggests a strong resilience to developing depression under typical circumstances. However, maintaining a healthy lifestyle, including regular physical activity, a balanced diet, and stress management practices, can further support mental well-being

# **Mood-induced Sleeplessness Risk**



**Neutral** 

**Very Low** 

### **Understanding what this may mean**

Individuals with this result are likely to have a neutral risk of developing sleeplessness as a result of mood or emotional fatigue. In general, low mood and emotional fatigue can interfere with sleep quality and lead to difficulties falling or staying asleep. However, individuals with this result are not at an increased risk compared to the general population.

# **Genetic Summary** 17

SNP Marker Genotype SNP Marker Genotype

| FKBP5 T106-2636C | rs1360780  | Indeterminate |
|------------------|------------|---------------|
| FKBP5 C*1136A    | rs3800373  | AA            |
| SOD2 A47G        | rs4880     | AG            |
| ACE A-239T       | rs4291     | AA            |
| MTHFR A359+160G  | rs17367504 | AA            |
| ESR1 A453-351G   | rs9340799  | AA            |
| DIO2 C-143T      | rs12885300 | TT            |
| DIO2 T274C       | rs225014   | TT            |
| NR3C1 G1184+646C | rs41423247 | CG            |

| CRHR1 C241+1631T  | rs242939   | TT |
|-------------------|------------|----|
| BDNF C196T        | rs6265     | CC |
| ARNTL T671-305G   | rs11022778 | TT |
| ABCC9 G3474-1222A | rs11046205 | GG |
| CRHR1 A122-1309C  | rs242941   | CC |
| TPH2 g.G4298T     | rs4570625  | TT |
| CNR1 C1359T       | rs1049353  | TC |
| FAAH C385A        | rs324420   | CC |