

STRESS RESISTANCE ON THE EXAMPLE OF SUPRAMOLECULAR-GENETIC LEVEL OF PLANT DEVELOPMENT

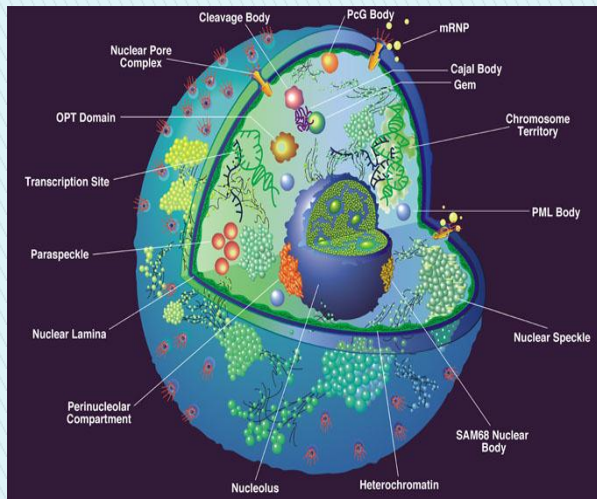
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(Ideological direction of the epigenetic phenotype (from Doctor of Biological Sciences R.N. Churaev)
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From the standpoint of the concept of **supramolecular** biochemistry (non-covalent interactions), on a model object of the epigenetic system of transitions of a spring wheat cultivar to a stress-resistant winter cultivar, an analysis of the barrier localization of proteomic *super*molecular assemblies on the interface of the cell nucleus was carried out: on the **suprablocks** - nucleoplasma (Np), eu- (Chr-I), heterochromatin Chr-II), nuclear matrix (NM) of the total chromatin matrix (TChrM).

Biochemical analysis of cell nuclei induced to organ-specific, coordinated-regular growth, hatching wheat germ

Cell nucleus



1.1. Age of hatching embryos, h (collection seeds from VIR)					
24	30	36	42	organs	48
Whole				Coleoptile Mesocotyl Root	
High-differential				Embryo-germ	
1.2. Isolation of cell nuclei					
1.3 Isolation of supra -structure-blocks with an increase in the ionic strength of the salt gradient that helps to weaken the electrostatic interaction					
0,14 M NaCl	0,35 M NaCl	2M NaCl	6M GuHCl		
Nucleoplasm, Labile chromatin	"Eu" chromatin loosely bound to NM	"Hetero" chromatin tightly bound to NM	Nuclear matrix		
(Np)	(Chr-I)	(Chr-II)	(NM)		
1.4. Gradient elution with GuHCl-guanidine hydrochloride, at the interface between non-histone and histone-supercomplex-ensembles of supra-structure-blocks, by ion exchange chromatography on IRC-50 (Heidelberg), prepared for work according to the description [Ivanova et al., 1975]					
Step concentrations					
6,0 - non-histone acidic proteins - (Ngb)					
8,9 - lysine-rich, linker histone - (HI)					
10,6- moderately lysine-rich histones - (H2A+H2B)					
13,0 - arginine-fortified histones (H3+H4)					
40,0- arginine-rich histones - (H3+H4)"					

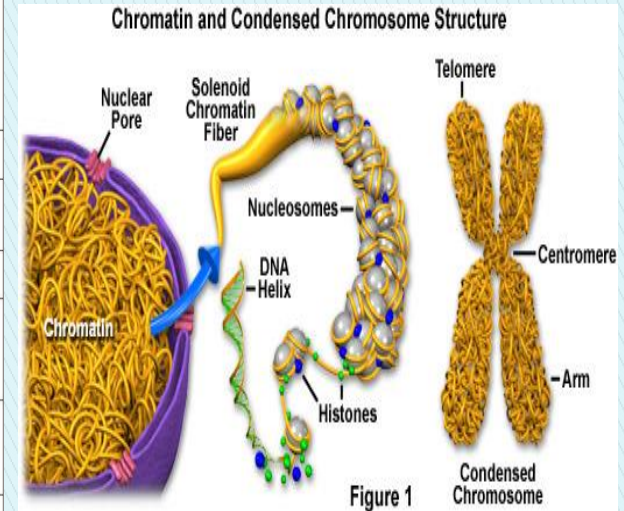
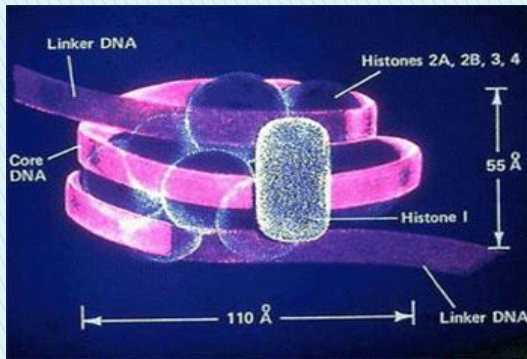


Figure 1

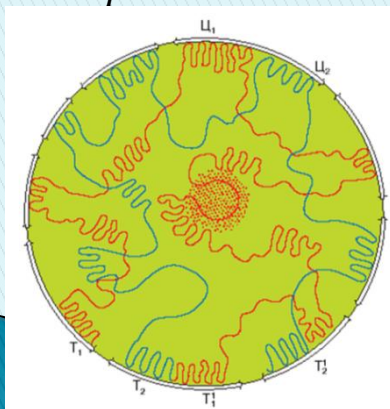
Features of the reorganization of the *super*molecular proteome at the interface of topologically associated supra-blocks of TChrM in the process of initiation of growth morphogenesis of the genetic subsystems of the whole organism of spring and winter wheat derived from it

Nucleosome

Mathematical Model

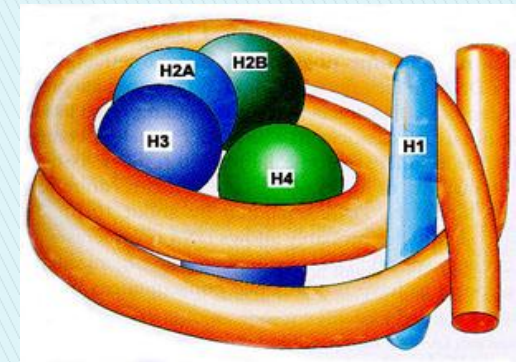


Arrangement of chromosomes in the interphase nucleus

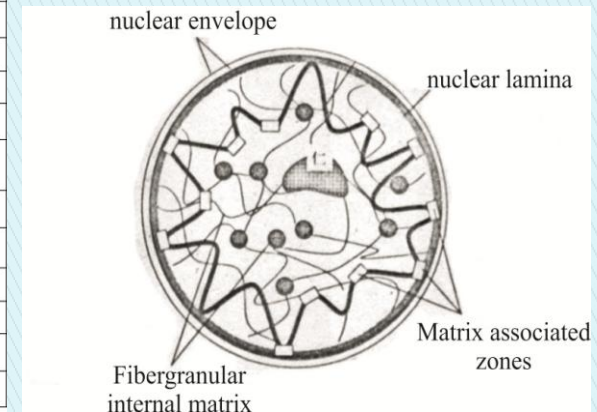


Total chromatin blocks	Organogenesis	Proteomic <i>super</i> molecular assemblies-structures	
		Spring	Winter
42 ч.			
Np	Coleoptile	HI	$(H2A+H2B) \geq HI \geq Hr6$
	Mesocotyl	$(H3+H4)''$	$HI \geq Hr6$
	Root	HI	$(H3+H4)'' \geq (H3+H4)'$
Chr-I	Coleoptile	$(H2A+H2B)$	$(H3+H4)'$
	Mesocotyl	$(H3+H4)''$	$(H2A+H2B)$
	Root	$(H2A+H2B)$	$(H2A+H2B) \geq HI$
Chr-II	Coleoptile	$(H3+H4)'$	$HI \geq Hr6$
	Mesocotyl	HI	$(H3+H4)'$
	Root	HI	$(H3+H4)'+(H3+H4)''$
NM	Coleoptile	$(H2A+H2B)$	$(H3+H4)'$
	Mesocotyl	Hr6	Hr6
	Root	$Hr6 \geq HI$	$Hr6 \geq HI \geq (H3+H4)''$
48 ч.			
Np	Coleoptile	$(H3+H4)'$	Hr6
	Mesocotyl	$(H2A+H2B)$	HI
	Root	HI	$(H2A+H2B) \geq Hr6$
Chr-I	Coleoptile	Hr6	$HI \geq (H3+H4) \geq Hr6$
	Mesocotyl	$Hr6 \geq (H3+H4)'$	$(H3+H4)'$
	Root	$(H2A+H2B) \geq HI \geq H3+H4)'$	$(H3+H4)'$
Chr-II	Coleoptile	$(H2A+H2B)$	$HI \geq (H3+H4)'$
	Mesocotyl	$(H3+H4)'$	$(H2A+H2B) \geq (H3+H4)'$
	Root	$(H3+H4)'$	$HI \geq Hr6$
NM	Coleoptile	$HI \geq Hr6$	Hr6
	Mesocotyl	$(H3+H4)'$	$Hr6 \geq HI \geq (H2A+H2B)$
	Root	$(H2A+H2B)$	HI

Core histone model

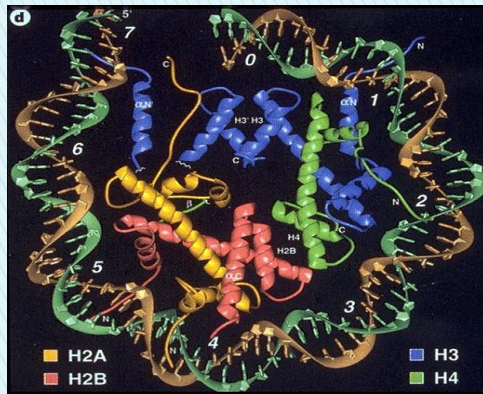


Arrangement of fibrogranular and fibrillar structures in the cell nucleus

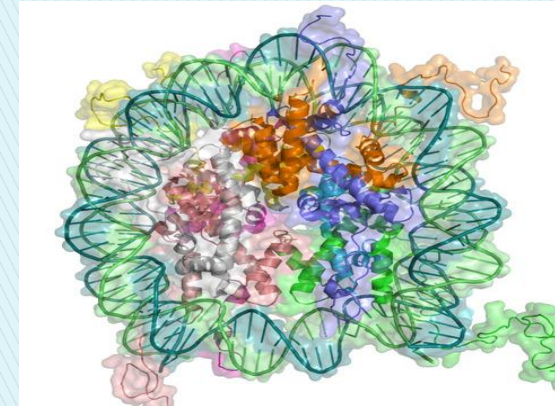


Supramolecular-genetic derivation

Nucleosome molecular model



An algorithm for the dynamics of proteomic *super*molecular ensembles on the interface of **supramolecular** blocks of TChrM is shown, on which (**Np, Chr-II, NM**) the positioning role of core (**H3 + H4**)" histones in the winter phenotype in the zone of geomorphism of the root system during **42** hours of integration stabilization of physiological genetic stress resistance of the organism. Thus, the proteomic genomics algorithm of integrative physiology leads to the concept of the perspective of the epigenetic landscape of the polyhedron.



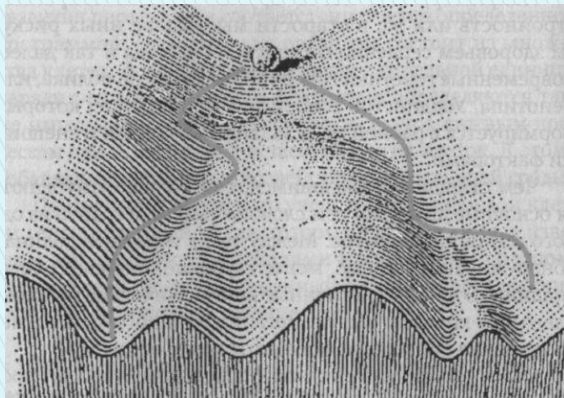


The motives of the epigenetic landscape in the depiction of artists

British scientist, born in India Waddington, Conrad Hal 08.11.1905–26.09.1975



Knight at the crossroads. Victor Mikhailovich Vasnetsov



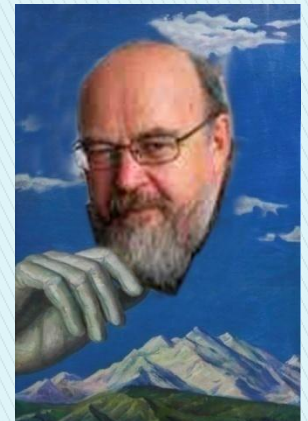
The epigenetic landscape of Waddington



Memories of Baikal



Overcoming



Reflections from above

Nikas Stepanovich Safronov