

**® 16. kongres Světové  
společnosti pro nervosvalová  
onemocnění (WMS),  
Portugalsko, říjen 2011**

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**Klinika dětské neurologie,  
MU a FN Brno**

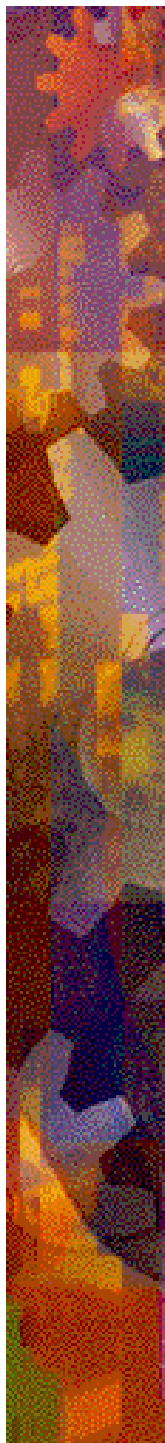



**16<sup>th</sup>** International Congress of the  
World Muscle Society

**Petr Vondracek**

Czech Republic

**October 18<sup>th</sup> - 22<sup>nd</sup>, 2011**  
Almancil | Algarve | Portugal





**Světový kongres s hlavním tématem:  
Léčba dědičných nervosvalových  
chorob, zejm. DMD/BMD a SMA**

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Parent Projectu**



## 1) Naše aktivní účast

® Posterová prezentace

® „Spectrum of point mutations in Czech DMD/BMD patients and their phenotypic outcome“

# Spectrum of point mutations in Czech DMD/BMD patients and their phenotypic outcome.



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## Introduction:

Duchenne and Becker muscular dystrophies (DMD/BMD) are X-linked recessive neuromuscular diseases caused by a mutation in the *DMD* gene. 60% of DMD/BMD cases are caused by deletions, 5% by large duplications, and 35% by point mutations. DMD is typically connected with mutations causing a premature termination codon (PTC), while BMD with in-frame del

## Methods:

- Mutation screening on mRNA level: reverse transcription-PCR, protein truncation test, DNA sequencing.
- Mutation screening on DNA level: PCR-sequencing.
- Muscle biopsy + immunohistochemistry: dystrophin (DYS1, 2, 3), N-trophin

## Czech DMD/BMD patients with the mutation in the *DMD* gene

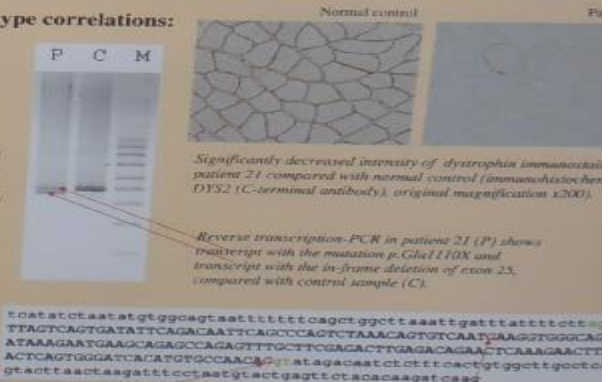
Patient	Phenotype	Location of mutation (exon)	Mutation identified on mRNA level	Mutation identified on DNA level	Mutation at protein level	Immunohistochemical labeling using antibodies DYS1,2,3
1	BMD	3	c.163G>T, 94_649del	not performed	p.Glu55X, P32_D217del	not performed
2	BMD	intron 3	c.187_264del	c.187-13_194ins21	p.Pro63_Asn88del	decreased intensity of DYS1 and DYS3
3	DMD	intron 4	c.264_265ins265-295_268-464	c.265-463A>G	p.Val89Met8X10	negative
4	DMD	7	c.303C>T	not performed	p.Arg195X	negative
5	DMD	7	c.303C>T	not performed	p.Arg195X	negative
6	DMD	intron 7	c.531_649del	c.649-5G>A	p.Pro178Cys6X2	DYS2 and DYS3 negative, DYS1 present at 20% of muscle fibres
7	DMD	intron 7	not performed	c.649-5G>A	p.Pro178Cys6X2	not performed
8	DMD	8	c.773C>T	not performed	p.Glu241X	not performed
9	DMD	9	c.806G>T	c.836G>T	p.Gly285X	negative
10	DMD	10	c.1062C>A	not performed	p.Trp354X	negative
11	DMD	11	c.1334C>T	not performed	p.Glu442X	negative
12	BMD	intron 12	c.1483_1662del	c.1483-4T>G	p.Val495_Lys534del	DYS1 negative, DYS2 and DYS3 decreased intensity
13	DMD	16	c.1873C>T	not performed	p.Glu625X	negative
14	DMD	16	c.1961T>G	c.1961T>C	p.Leu548X	not performed
15	DMD	18	c.2276T>G	not performed	p.Leu759X	negative
16	DMD	18	c.2281G>T	not performed	p.Glu761X	negative
17	DMD	20	not performed	c.2566C>T	p.Glu856X	not performed
18	DMD	intron 20	c.2622_2623insAG	c.2623-3C>G	p.Asp875Asp6X17	negative
19	DMD	21	c.2797C>T	not performed	p.Glu933X	negative
20	BMD	24	c.[...3185_3239del]	c.3236-113>A	p.Asn1093_Arg1092del p.His1118X, Leu1093_Glu1144del	decreased intensity
21	BMD	25	c.3328G>T, 3277_3435del	not performed	p.Val495_Lys534del	DYS1 and DYS2 normal, DYS3 decreased intensity
22	BMD	25	c.3277_3435del	c.3432G>A	p.Leu1093_Glu1144del	normal
23	BMD	intron 25	c.[...3777_3435del]	c.3432-1G>A	p.[...Leu1093_Glu1144del]	normal
24	BMD	intron 25	c.[...3277_3435del]	c.3432-2T>C	p[...Leu1093_Glu1144del]	DYS3 negative, DYS1 and DYS2 decreased intensity
25	DMD + CMT1A	27	c.3689_3613del TAAAAGCCTT	not performed	p.Lys1264Ile6X11	almost negative
26	DMD	28	c.[3823C>G, 3797_3971del, 3792_3935del]	c.3823C>G	p.Trp1774X, Glu1263_Asp1307del, Glu1263_Glu1376del	normal
27	DMD	29	not performed	c.3962C>T	p.Glu1378X	not performed
28	DMD	30	c.4099C>T	not performed	p.Glu1367X	negative
29	DMD	33	c.4636delA	not performed	p.Ala1547Leu6X2	negative
30	BMD	intron 33	c.[...4674_4675ins18]	c.4675-11A>G	p.[...Val1589Phe6X28]	DYS1 negative, DYS2 and DYS3 decreased intensity
31	DMD	34	c.4729C>T	not performed	p.Arg1577X	negative
32	DMD	36	c.5011C>A	not performed	p.Tyr1677X	negative
33	DMD	39	c.5474C>T	not performed	p.Glu1826X	negative
34	DMD	39	c.5583delC	not performed	p.Glu1835Asp6X13	negative
35	DMD	40	c.5853C>T	not performed	p.Glu1893X	negative
36	DMD	40	c.5662delC	not performed	Asp1888Ile6X3	negative
37	DMD	42	c.5967C>G	not performed		

## Results:

- Excluding deletions and duplications by MLPA (Multiplex Ligation dependent Probe Amplification), 65 different point mutations in 63 DMD/BMD patients and 5 DMD carriers were identified. 38 types of mutations were unique for Czech DMD/BMD population (mentioned by bold letters in the Table).
- 51 patients have DMD phenotype, 12 patients BMD phenotype.
- Two patients (3 and 55) have point mutations localised deep in introns. These mutations create new splicing signals and cause insertions of a part of intron sequence into mRNA.
- One patient (49) carries an insertion of the repetitive sequence *AluYa5*. This insertion disrupted the splicing signal and led to the frame-shift deletion.
- Four patients have mutations creating PTC and BMD phenotype (patients 1, 21, 26 and 27). All of these cases can be explained by molecular mechanisms connected with expression of genetic information (dystrophin production is initiated at the new start codon, mutations change an splicing enhancer sequence and cause in-frame deletions, mutations near the 3' end can escape from mediated mRNA decay).

## Cases with interesting genotype-phenotype correlations:

**Patient 21** – the mutation c.3328G>T, p.Glu1110X; BMD phenotype – the mutation changes an exon splicing enhancer sequence in exon 25 and causes in-frame deletion of exon 25 (predicted by Rescue-ESE program, <http://genes.mit.edu/burgelab/rescue-ese/>).



Significantly decreased intensity of dystrophin immunoblotting in patient 21 compared with normal control (immunohistochemical DYS2 (C-terminal antibody), original magnification x200).

Reverse transcription-PCR in patient 21 (P) shows transcript with the mutation p.Glu1110X and transcript with the in-frame deletion of exon 25, compared with control sample (C).


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c a a t a c t a a a t a t g t g g c a g t a a t t t t t t c a g c t a a a t t g g c t a a a t t g a t t a t t t t o t s ;
T T A G C C A G T G A T A T T C A G A C A A T T C A G C C C A G T C T A A C A C T C T C A A T C A A G G T G G G C A ;
A T A A A G A A T G A A G C A G A G C C A G A G T T T G C A C T T G A G C A G A C T C A A G A C T T
A C T C A G T G G G A T C A C A T G P G C C A A C G G T a t a g a c a a t c t d t t c a c t g t g g g t g o c t
g t a c t t a a c t a a g a t t t c c t a a t g t a c t g a g t t c t a c a a a g a t c a a g
    
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Point mutations in exon 25 (red letters) causing exon skipping in patient 21 and patient 22 (exon sequence is mentioned by capital letters, intron conserved sequences - by green letters).

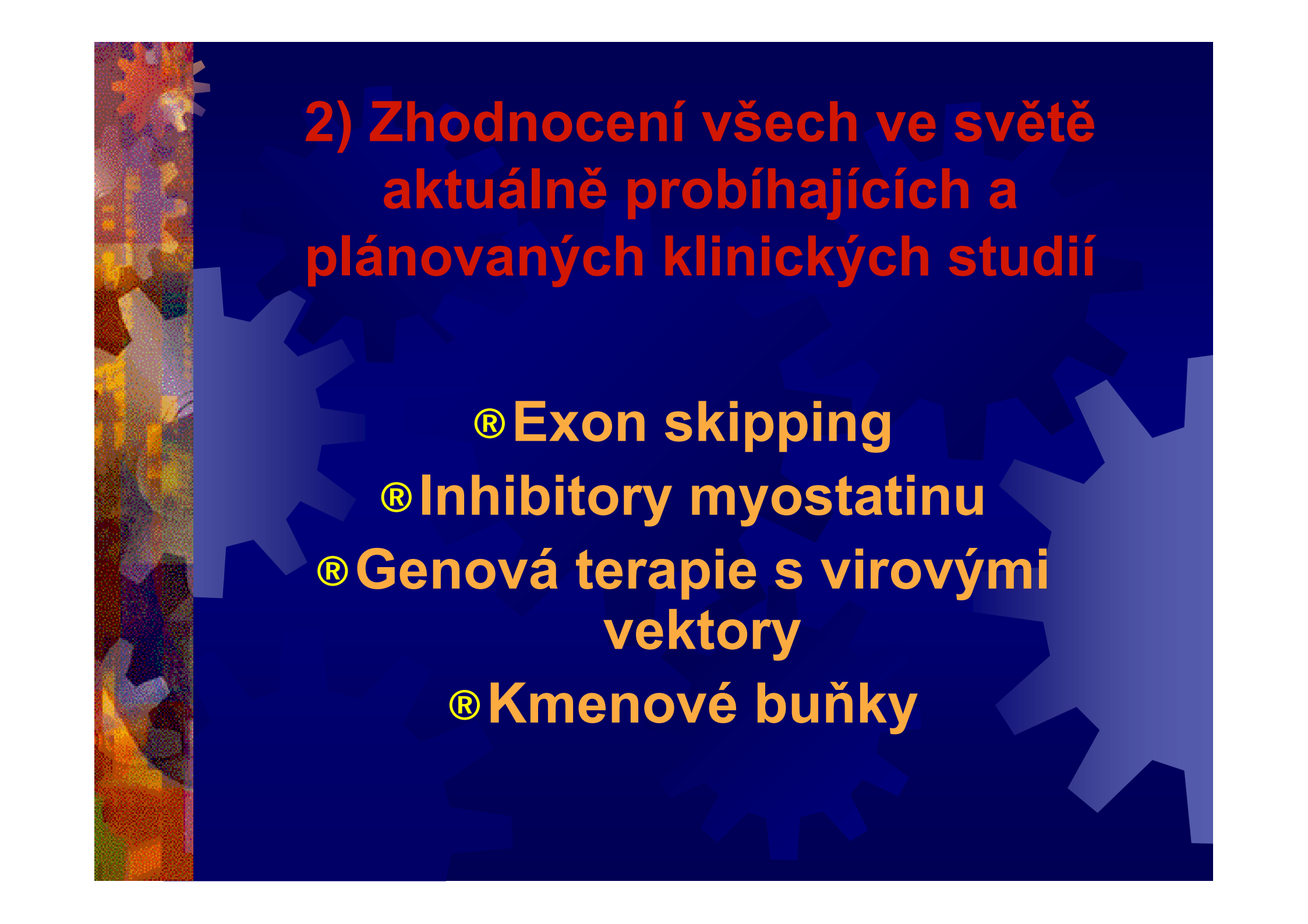
## Insertions of parts introns into mRNA caused by single nucleotide changes deep in introns:

- Patient 3** – the mutation c.265-463A>G caused creation of a novel splice site, and the insertion of
- Patient 55** – the mutation c.9564-427T>G caused creation of a novel splice site, and insertion of



**Práce prezentuje bodové mutace v DMD genu u všech 63 pacientů a 5 přenašeček DMD/BMD na podkladě bodových mutací, kteří byli dosud diagnostikováni v ČR**

- ® Význam pro budoucí molekulárně genetickou terapii DMD/BMD**
- ® Práce živě diskutována a velmi oceněna zejm. holandskými specialisty na problematiku DMD/BMD**



## 2) Zhodnocení všech ve světě aktuálně probíhajících a plánovaných klinických studií

® Exon skipping

® Inhibitory myostatinu

® Genová terapie s virovými  
vektory

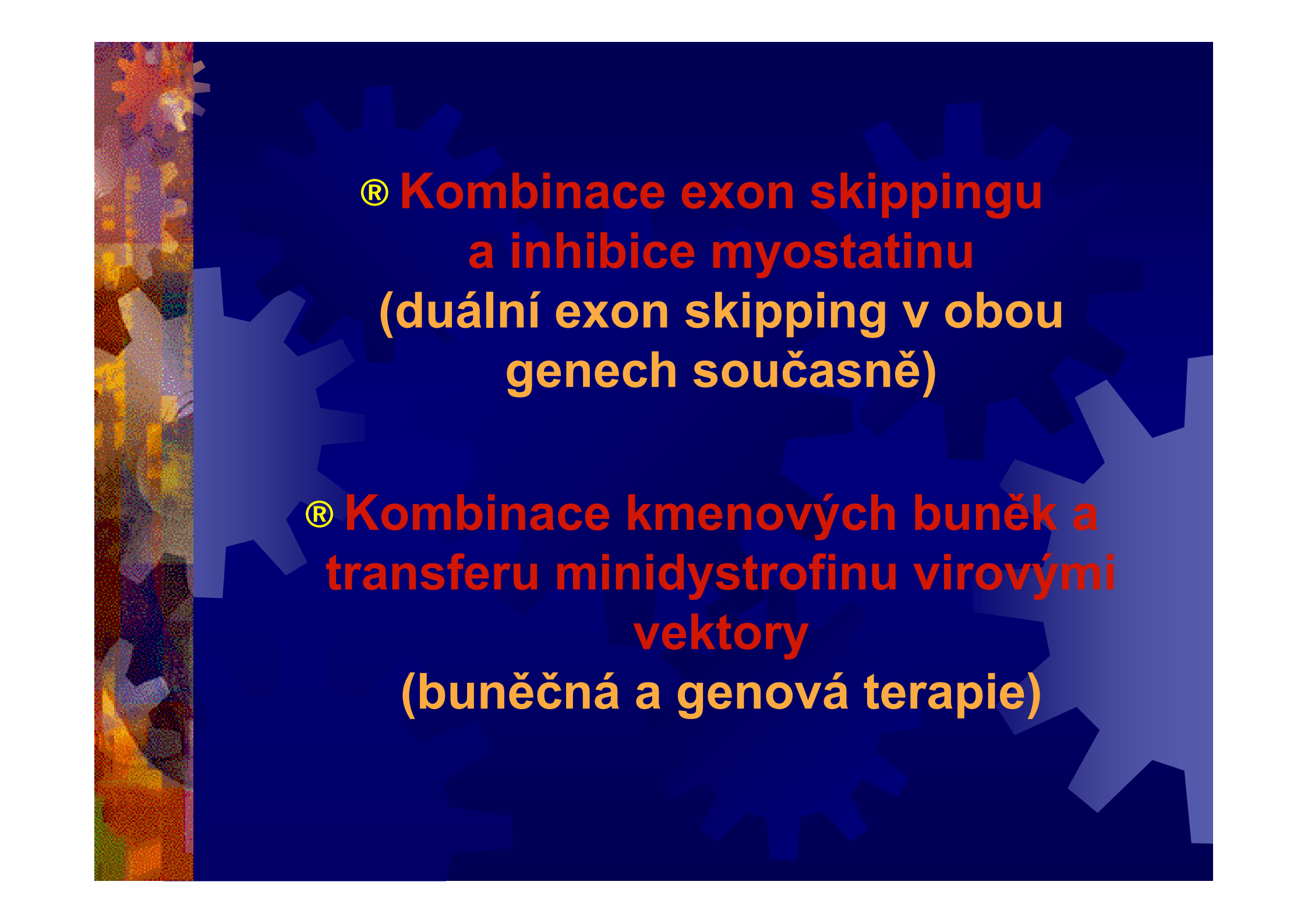
® Kmenové buňky

### 3) Horké novinky

® Kromě pokračujícího testování exon skippingu **51** ( III. fáze) a **44** ( I/II. fáze), již od 2012 klinické testování skippingu dalších exonů:  
**45 a 53** (1. polovina 2012)  
**52 a 55** (preklinický vývoj)

® Rozšíření spektra delecí + již také duplikací, které bude možno léčit exon skippingem






® **Kombinace exon skippingu  
a inhibice myostatinu**  
(duální exon skipping v obou  
genech současně)

® **Kombinace kmenových buněk a  
transferu minidystrofinu virovými  
vektory**  
(buněčná a genová terapie)

## 4) Osobní kontakty

- ® diskuse s hlavními koordinátory klinických studií ve světě a výměna informací o naší studii DMD114044 a dalším společně plánovaném a koordinovaném postupu při testování terapie DMD/BMD
  - ® Natalie Goemans (Leuven, Belgie)
  - ® Thomas Voit (Paříž, Francie)
  - ® Kate Bushby (Newcastle, UK)
  - ® Kevin Flanagan (Columbus, Ohio, USA)



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