

# MIBio 2015

## *Drug Discovery – Some Current Trends*

**Dr Alan E Smith CBE FRS**

*Chairman, Cambridge in America*

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October 21<sup>st</sup> 2015

Magdalene College  
Cambridge, UK

Drug discovery is hard

Rare diseases

'All diseases are rare'

Alternative approaches; immunotherapy

Current biotech bubble

Formulation

# Genetic Code, Initiation, & Eukaryotic Protein Synthesis

## LMB, Cambridge 1967-70

### Cytoplasmic Methionine Transfer RNAs from Eukaryotes

by  
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TRANSFER RNA from bacteria contains two methionine accepting species  $tRNA_{Met}^{bact}$  and  $tRNA_{Met}^{euc}$ . Methionine attached to  $tRNA_{Met}^{euc}$  can be enzymatically formylated to give formylmethionyl  $tRNA_{Met}$  (fMet- $tRNA_{Met}$ ), while methionyl  $tRNA_{Met}$  (Met- $tRNA_{Met}$ ) does not accept formyl groups<sup>1</sup>. fMet- $tRNA_{Met}$  is the chain initiator on the 70S ribosomes of all prokaryotes and prokaryotic-type chloroplasts and mitochondria<sup>2</sup>. A formylated initiator tRNA donates methionine to the amino group of newly synthesized polypeptides.

Eukaryotic cells contain two species of methionine accepting tRNA. One of these (Met- $tRNA_{Met}^{euc}$ ) only incorporates methionine at terminal positions of polypeptides whereas the other (Met- $tRNA_{Met}^{euc}$ ) donates methionine into internal positions. It is suggested that the latter is an initiator tRNA.

(Reprinted from Nature, Vol. 225, No. 5228, pp. 184-187, January 10, 1970)

### Translation of RNA from Encephalomyocarditis Virus in a Mammalian Cell-free System

THE role of formylmethionyl  $tRNA_t$  (fMet- $tRNA_t$ ) in the initiation of protein biosynthesis in *Escherichia coli* is now reasonably well understood<sup>1</sup>. A similar mechanism involving fMet- $tRNA_t$  seems to be common to all 70S type ribosomal systems including those from bacteria, chloroplasts and mitochondria<sup>2,3</sup>. The corresponding process on 80S type ribosomes is, however, as yet unknown, but there is no evidence to suggest that fMet- $tRNA_t$  or any similar blocked aminoacyl-tRNA is involved<sup>2</sup>. A mammalian cell-free system which syn-

### The size of Roundly active in cell-free

Tony Pawson<sup>\*</sup>, Robert Harvey &  
Department of Molecular Virology, Imperial Cancer Research Fund, London WC2, UK

Sedimentation analysis of mRNA from RSV-infected cells suggests that Pr76, the precursor to the viral RNA, is synthesised on a mRNA of similar size to the viral glycoprotein, gp85, is synthesised on a smaller size class of mRNA.

### Initiator Codons in Eukaryotes

by  
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THE cytoplasm of several eukaryotes contains two Methionine accepting tRNAs ( $tRNA_{Met}^{euc}$  and  $tRNA_{Met}^{bact}$ ). Methionine attached to  $tRNA_{Met}^{euc}$  can be formylated by *Escherichia coli* transformylase<sup>1</sup>, while Met- $tRNA_{Met}^{euc}$  does not accept formyl groups. In an *E. coli* cell-free system eukaryotic fMet- $tRNA_{Met}^{euc}$  functions as an initiator with the same requirements and specificity as *E. coli* fMet-

ROUS sarcoma virus (RSV) contains a genomic virion 70S RNA which in denaturing conditions dissociates into two apparently identical 30-40S RNA in cell-free systems have translated RSV RNA. The major *in vitro* product derived from mammalian cells<sup>1</sup>. We and others of RSV virion 30-40S RNA is a polypeptide similar, if not identical, to the 76,000 molecular weight precursor to group-specific (gs) antigens, Pr76, which is found in RSV-infected cells<sup>2</sup>. Pr76 synthesised *in vitro* in response to RSV virion 30-40S RNA can be labelled with N-formyl-methionine derived from the initiator tRNA<sub>Met</sub>. Since fMet can discriminate against the Met-tRNA<sub>Met</sub> by the corresponding aminoacyl-tRNA binding

in a different amount and translated at a different efficiency, the viral proteins need not be synthesised in equimolar amounts. This model also suggests that virion 30-40S RNA has internal initiation sites and that they are inactive. To examine the mechanism by which RNA tumour virus proteins are synthesised, we have isolated total poly(A)-containing cytoplasmic mRNA from RSV-infected chick cells. This mRNA has been fractionated on denaturing sucrose gradients, translated in mammalian cell-free systems derived from mouse ascites and L cells. Because infection by RNA tumour viruses does not shut off host-cell metabolism and viral mRNA constitutes only a small fraction of total mRNA activity, the viral proteins made *in vitro* were isolated by immunoprecipitation using various antisera raised against avian RNA tumour virus proteins.

### Fractionation of infected cell mRNA

Poly(A)-containing cytoplasmic RNA was isolated from e/o chick embryo fibroblasts transformed with non-defective Prague-C RNA, from uninfected chick cells, and from



# Function of Tumor Antigens and Identification of Nuclear Localisation - Mill Hill, London 1980-1984

Reprinted from Nature, Vol. 311, No. 5981, pp. 33-38, 6 September 1984  
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## Sequence requirements for nuclear location of simian virus 40 large-T antigen

Daniel Kalderon, William D. Richardson, Alexander F. Markham\* & Alan E. Smith

Cell, Vol. 18, 915-924, December 1979. Copyright © 1979 by MIT

## Protein Kinase Activity Associated with Polyoma Virus Middle T Antigen in Vitro

Alan E. Smith, Ros Smith, Ben Mike Fried†  
Translation, Nucleic Acid\* and Genetics‡ Laboratories  
Imperial Cancer Research Fund  
Lincoln's Inn Fields  
London WC2, England.

### Summary

A protein kinase activity can nonprecipitates of extracts from infected cells using antiserum transformed cells (anti-T serum) detected in uninfected cells serum. Using rat anti-T serum the heavy chain of rat IgG

Cell, Vol. 39, 499-500, December 1984 (Part 2). Copyright © 1984 by MIT

## A Short Amino Acid Sequence Able to Specify Nuclear Location

Daniel Kalderon, Bruce L. Roberts, William D. Richardson, and Alan E. Smith  
Biochemistry Division  
National Institute for Medical Research  
Mill Hill  
London NW7 1AA, England

### Summary

A short sequence of amino acids including Lys-128 is required for the normal nuclear accumulation of wild-type and deleted forms of SV40 large T antigen

## Polyoma virus transforming protein associates with the product of the c-src cellular gene

Sara A. Courtneidge & Alan E. Smith  
Biochemistry Division, National Institute for Medical Research,  
Mill Hill, London NW7 1AA, UK

Reprinted from Nature, Vol. 303, No. 5916, pp. 435-439, 2 June 1983  
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below M<sub>r</sub> 15,000 is not detectably retarded, a globular protein of M<sub>r</sub> 67,000 (BSA) enters nuclei only very slowly, and a protein of M<sub>r</sub> 450,000 (ferritin) is excluded (Paine et al., 1975). From such studies, the nuclear envelope has been proposed to behave as an array of cylindrical pores of internal diameter 7-11 nm (Paine et al., 1975; Peters, 1984). This model of the envelope corresponds reasonably well with the dimensions of discrete structures called nuclear pores, which have been purified from the nuclear envelope and analyzed by electron microscopy (Harris, 1981; Urwin and Milligan, 1982).

It appears therefore that in vivo many small proteins may diffuse freely between nucleus and cytoplasm and may

is good evidence that it has an intrinsic kinase activity that is directly involved in RSV transcription. Kinase activities of pp60<sup>src</sup> or its cellular homologues are assayed in immunoprecipitates of serum rabbits bearing RSV-induced tumours (anti-T serum) transferred to tyrosine in two alternative systems: immunoglobulin<sup>1,2</sup> or middle T<sup>3,4</sup> or both<sup>5,6</sup>. When extracts from polyoma virus-transformed cells are assayed in the same way using serum polyoma virus-induced tumours (anti-T serum) observed. Using hamster anti-T serum, middle T<sup>3,4</sup> or middle T<sup>5,6</sup> or both<sup>7,8</sup> or both<sup>9,10</sup> are capable of autophosphorylation can be detected in the TBR antibody directed immunoprecipitates of either normal or transformed rat cells were formed from polyoma virus-transformed rat cells were formed alone, middle T<sup>3,4</sup> or middle T<sup>5,6</sup> was still





## Drug Discovery is hard

We still have much to learn about human biology and disease

Takes a long time; costs a great deal; most things fail

For last 30+ years, as more money has been spent in pharma R&D, less drugs have been approved

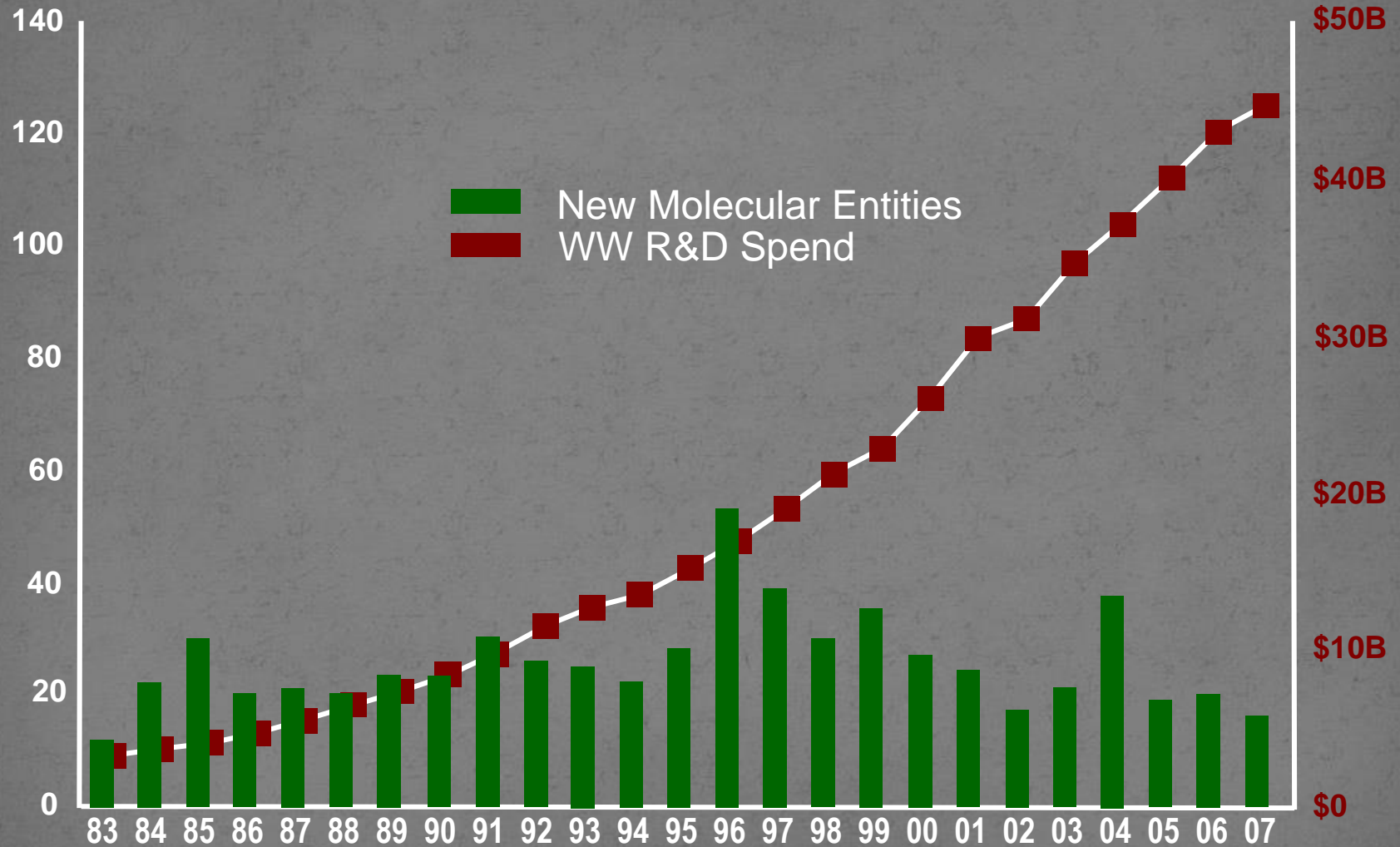
Most pharma have concluded this is a poor investment and that they are unable to discover drugs in house

Current model is to outsource much early research

Pharma become Development and Sales & Marketing

*Does outsourcing discovery work any better?*

For 25 years Biopharma R&D spending grew steadily ...  
but paradoxically no correlation with more novel drugs approved



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## Opportunity for start ups

Innovation greater in early start ups?

Costs in start ups much less; Infrastructure minimized

New ideas pushed rapidly to proof of concept in early human trials

Most of these also fail

Successes valued hugely

*Overall do the finances work better in this model?*

## Biotech Madness – a recent example

Alexion acquires Synageva for \$8.4 bn - Spring 2015

Only drug: Kanuma (sebelipase alfa), a treatment for the ultra rare lysosomal acid lipase deficiency (LAL-D)

Recently approved EU, presently under FDA's priority review program,

Alexion made its name with Soliris, another ultra rare disease treatment (\$500,000 a year)

*This valuation is crazy*



## Rare Orphan diseases

Considered to be more likely to succeed

Molecular mechanisms established, in humans

Therapy often replacement, rather than interventional

Few patients mean that price is very high

Value created by high efficacy in most patients

Pioneered by Genzyme, now universal

*As more developed, total cost becoming prohibitive?  
(1% of drugs cost 30% of total spending)*

## All diseases are rare

As molecular mechanisms of disease unraveled  
many major diseases sub-categorised

Each sub category becomes 'rare'

Example, breast cancer; at least a dozen mechanisms

Rare disease approach yields higher success rates

Price for each drug necessarily will be very high

*Total drug costs will become unsustainable*



## New needs and opportunities

A general approach to cancer is very attractive

Immune involvement in control of cancer known for 100+ years – Colley's adjuvant

Earlier approaches unsuccessful – tumours developed resistance to immune attack

New approaches that circumvent immune counter attack (eg check point inhibitors) extremely promising

Early approvals in hand, multiple major programs

*A very promising 'novel' approach to cancer*

## New financial models

Early research and late development relatively easy to finance

Translational work, 'valley of death' more difficult

New models constantly tested

US Cystic Fibrosis Foundation recently sold Vertex royalty stream for \$3bn +

Foundations, patient organisations & hospitals will copy

*A new funding source with strong vested interest*



## Biotech Bubble

For several years biotech start up valuations have been extremely high

More recently in EU & UK

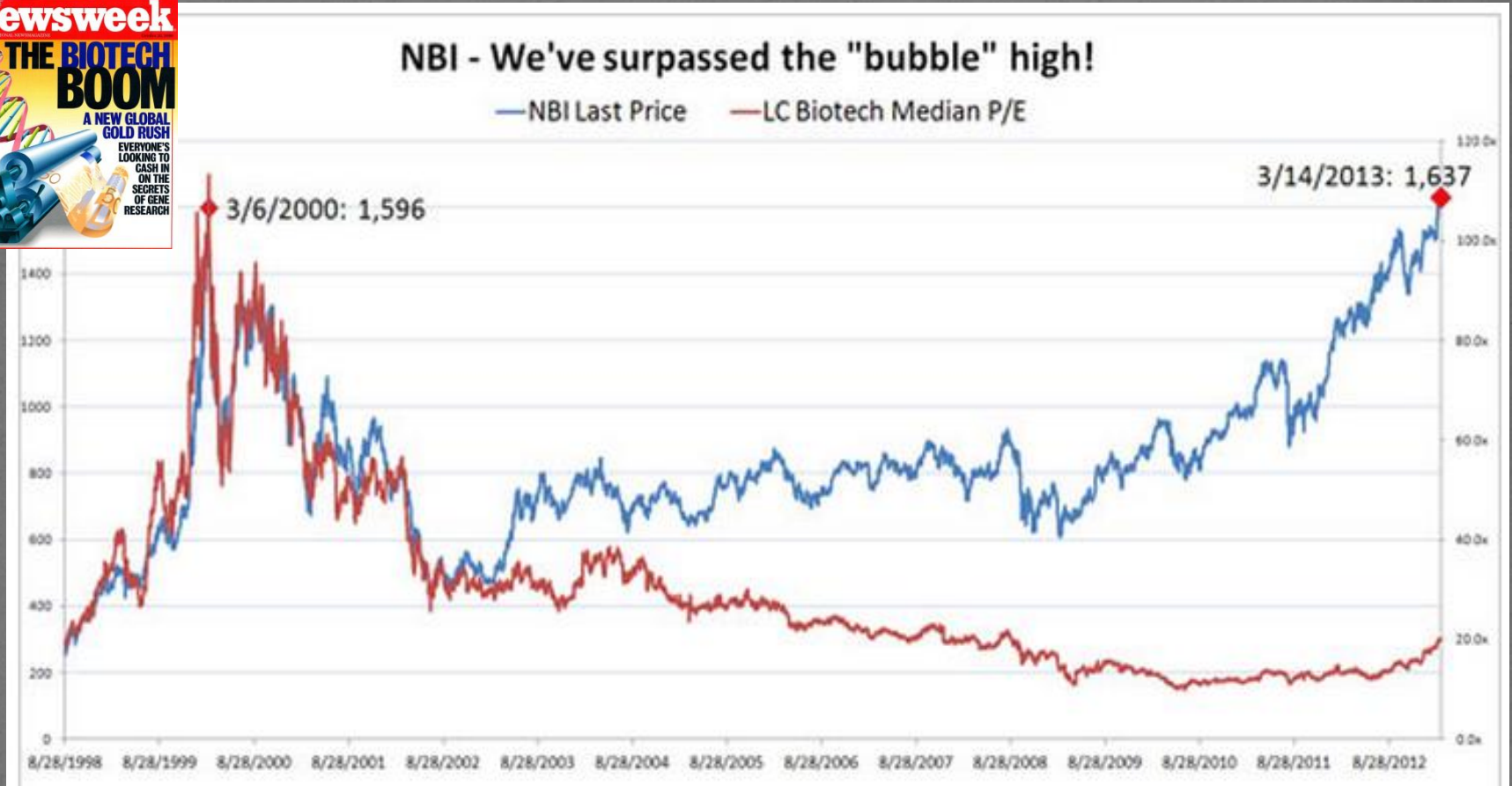
Biotech stocks outperforming for 5 years in US

Although turnover is high, the industry is thriving

This will not last

*Opportunism and luck are key components  
go for it while you can!*

# Q1 2013 ... Biotech Stocks Pass 2000 Peak Levels ...

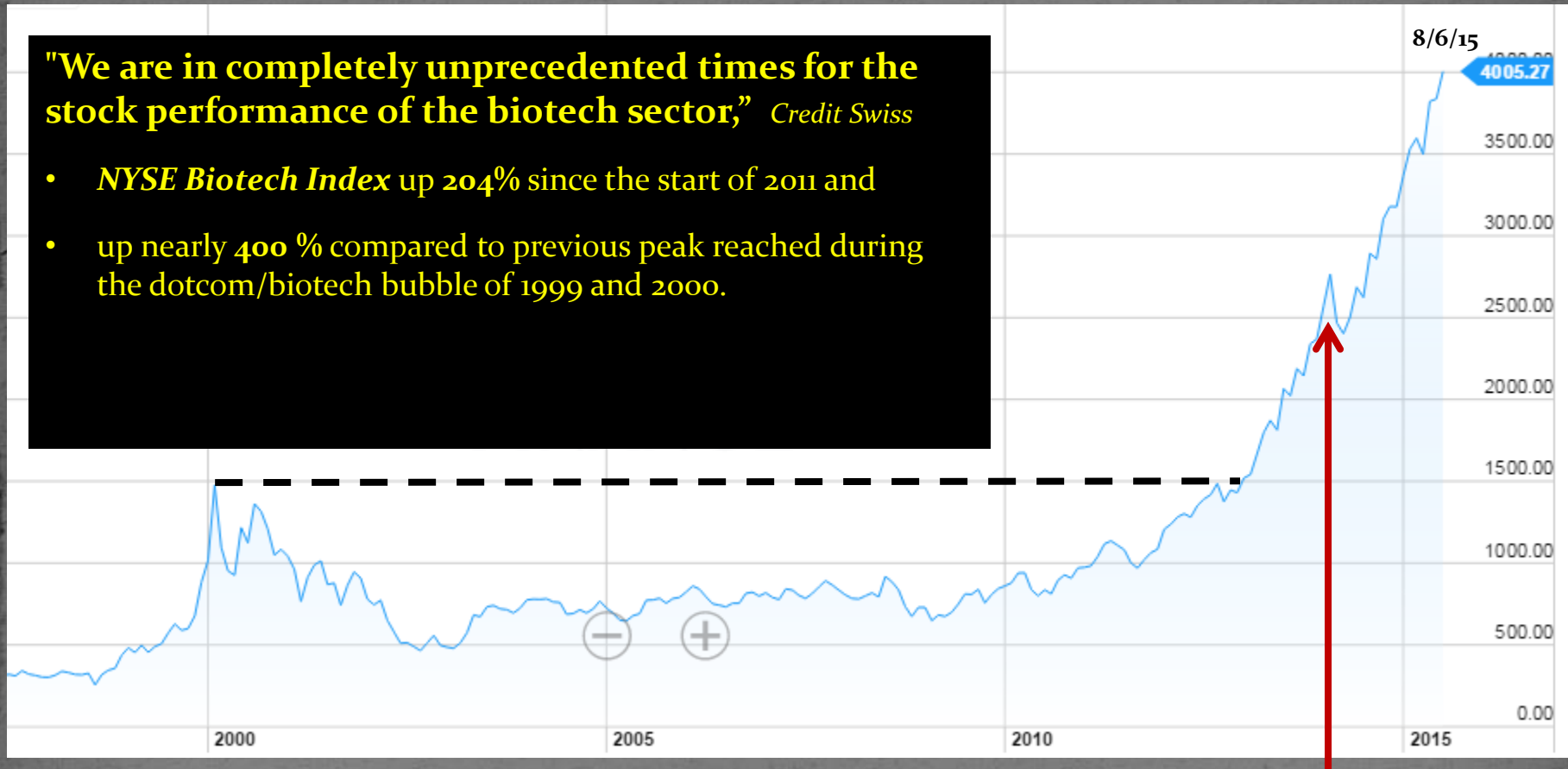


NBI = NASDAQ Biotech Stock Index

... and have kept on going!

"We are in completely unprecedented times for the stock performance of the biotech sector," *Credit Swiss*

- *NYSE Biotech Index* up 204% since the start of 2011 and
- up nearly 400 % compared to previous peak reached during the dotcom/biotech bubble of 1999 and 2000.



*Federal Reserve Chair Janet Yellen:*

... Some valuation metrics are substantially stretched, particularly Biotech



## Formulation

New drugs are relatively few

Add maximum value to existing proven drugs

Formulation can be key; particularly in biologics

Insulin has been a successful drug almost 100 years

Total sales insulin today higher than ever

*Formulation of existing drugs offers a low risk opportunity to add value for patients and investors*

Thank you

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